STIC-Biotech/ChemLib

162242

From:

Bowman, Amy

Sent:

Thursday, August 11, 2005 11:46 AM STIC-Biotech/ChemLib

To:

Cc: Subject: Bowman, Amy sequence search-10/665,951

I need SEQ ID NO: 2185 searched in application 10/665,951, length limited to 30 nucleotides.

Thank you, Amy Bowman AU 1635 **REM 2C31** mail REM 2C18 571-272-0755

somuch that

STAFF USE ONLY Searcher:_ Searcher Phone: 2-Date Searcher Picked up: Date Completed: Searcher Prep/Rev. Time: Online Time:_

Type of Search NA#: AA#: Interference:____ SPDI S/L:____ Oligomer:_ __ SPDI:_ S/L:_ Encode/Transl:_ Structure#:___ Inventor: Litigation:

******* Vendors and cost where applicable STN:_ DIALOG: QUESTEL/ORBIT: LEXIS/NEXIS:_ SEQUENCE SYSTEM:_ WWW/Internet:_ Other(Specify):_

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US-10-944-611-2809/c
                                                                                                                               August 18, 2005, 22:19:12 ; Search time 1735 Seconds (without alignments) 159.172 Million cell updates/sec
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/ntodata/1/pna/US10_NEW_COMB.seq7:*
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1: /cgn2 6/ptodata/1/pna/PCT_NEW_COMB.seq:*

2: /cgn2 6/ptodata/1/pna/PCT_NEW_COMB.seq:*

3: /cgn2 6/ptodata/1/pna/USO6 NEW_COMB.seq:*

4: /cgn2 6/ptodata/1/pna/USO8 NEW_COMB.seq:*

5: /cgn2 6/ptodata/1/pna/USO8 NEW_COMB.seq:*

6: /cgn2 6/ptodata/1/pna/USO9 NEW_COMB.seq:*

7: /cgn2 6/ptodata/1/pna/USO9 NEW_COMB.seq:*

8: /cgn2 6/ptodata/1/pna/USO9 NEW_COMB.seq:*

9: /cgn2 6/ptodata/1/pna/USO9 NEW_COMB.seq:*

10: /cgn2 6/ptodata/1/pna/USO0 NEW_COMB.seq:*

11: /cgn2 6/ptodata/1/pna/USI0 NEW_COMB.seq:*

12: /cgn2 6/ptodata/1/pna/USI0 NEW_COMB.seq:*

12: /cgn2 6/ptodata/1/pna/USI0 NEW_COMB.seq:*

13: /cgn2 6/ptodata/1/pna/USI0 NEW_COMB.seq:*
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GenCore version 5.1.6
Copyright (c) 1993 - 2005 Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                                                                          23768202 segs, 6575307184 residues
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Maximum Match 100%
Listing first 45 summaries
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Perfect score:
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

	;	Ap	Ap	Αp	Ap	Ap	Αp	Ap	ΑĎ
	_	2809,	2967,				3143,		2809,
	Description	Sequence	Seguence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence
SUMMARIES	ID	US-10-944-611-2809	US-10-944-611-2967	US-10-944-611-2977	US-10-944-611-3139	US-10-944-611-3141	US-10-944-611-3143	US-10-944-611-3144	US-10-962-898-2809
	DB	13	13	13	13	13	13	13	14
	Length	19	19	19	19	19	19	19	19
de	Query Match Length DB ID	90.5	90.5	90.5	90.5	90.5	90.5	90.5	90.5
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Sequence 2967, Ap	Sequence 2911, Ap	3141	3143,	3144,	296	N	296	29	(7	24, App	e 2742,	2745,	2763,	2765,	2767,	2769,	2777,	2778,	2843,	Sequence 2844, Ap	2845,	2846,	2864,	2865,	e 2868,	e 2872,	e 2876,	e 2880,	e 2925,	e 2928,	e 2931,	e 2934,	e 2937,	e 2940,	<u>.</u> `
10-962-898-2967	10-962-696-297	10-962-696-3139	10-962-898-3143	10-962-898-3144	10-944-611-2966	-944-611-2976	-962-898-2966	10-962-898-2976)-727-780A-23 S)-727-780A-24 S	10-944-611-2742	10-944-611-2745	10-944-611-2763	10-944-611-2765	10-944-611-2767	10-944-611-2769	10-944-611-2777	10-944-611-2778	10-944-611-2843	10-944-611-2844	10-944-611-2845	10-944-611-2846	10-944-611-2864	10-944-611-2865	10-944-611-2868	10-944-611-2872	10-944-611-2876	10-944-611-2880	10-944-611-2925	10-944-611-2928	10-944-611-2931	10-944-611-2934	10-944-611-2937	10-944-611-2940	10-944-611-2953
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ALIGNMENTS

US-10-944-611-2809/ Sequence 2809, Application US/10944611
Sequence 2809, Application US/10801
Sequence 2809, Application US/10801
Septicant: Jadhav, Vasant
APPLICANT: Sossen, Karl
Septicant: Vaish, Narendra
APPLICANT: Wisher Narendra
APPLICANT: Wisher Now Increment Endothelial Growth Factor And Vascular Endothelial Growth Factor Applicanton NUMBER: US/10/944,611
CURRENT APPLICATION NUMBER: US/10/944,611
CURRENT FILING DATE: 2004-09-16
PRIOR PILING DATE: 2004-04-23
PRIOR PELING DATE: 2003-09-16
PRIOR FILING DATE: 2003-09-16
PRIOR FILING DATE: 2003-09-16
PRIOR FILING DATE: 2003-09-16
PRIOR FILING DATE: 2003-09-16
PRIOR PELING DATE: 2003-09-16
PRIOR PELING DATE: 2003-09-16
PRIOR APPLICATION NUMBER: US 10/664,767
PRIOR APPLICATION NUMBER: US 10/605,255
PRIOR PELING DATE: 2003-09-16
PRIOR APPLICATION NUMBER: US 60/393,796

Gaps

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APPLICANT: Kossen, Karl
APPLICANT: Zinnen, Shawn
APPLICANT: Zinnen, Shawn
APPLICANT: Vaish, Narendra
APPLICANT: Waish, Narendra
APPLICANT: McSwiggen, James
TITLE OF INVENTION: RNA Interference Mediated Inhibition Of Vascular Endothelial
TITLE OF INVENTION: Growth Factor And Vascular Endothelial Growth Factor Receptor
TITLE OF INVENTION: Gene Expression Using Short Interfering Nucleic Acid (SiNA)
FILE REFERENCE: 400/235 (MBHB02-742-S)
CURRENT APPLICATION NUMBER: US/10/944,611
CURRENT FILING DATE: 2004-09-16
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US-10-944-611-2977
                                                                                                                                           siNA sense region
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PRIOR APPLICATION NUMBER: US 10/844,076

PRIOR PELING DATE: 2004-05-11

PRIOR PELING DATE: 2004-05-13

PRIOR PELING DATE: 2004-04-23

PRIOR PELING DATE: 2004-01-26

PRIOR PELING DATE: 2003-09-23

PRIOR PELING DATE: 2003-09-23

PRIOR PELING DATE: 2003-09-16

PRIOR PELING DATE: 2003-09-20

PRIOR PELING DATE: 2003-09-16

PRIOR PELING DATE: 2003-09-16

PRIOR PELING DATE: 2003-09-16

PRIOR PELING DATE: 2003-09-16

PRIOR PELING DATE: 2003-09-20

PRIOR PELING DATE: 2004-05-29

PRIOR PELING DATE: 2004-05-24

PRIOR PELING DATE: 2004-05-24
                                                                                                                                                                     FEATURE:
NAME/KEY: misc_feature
LOCATION: (1)...(1)
OTHER INFORMATION: 5'-3 attached terminal deoxyabasic moeity
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                                                                                                                                                                                                                                                                                                                                                                                          Length 19;
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                                                                                                   FEATURE: OTHER INFORMATION: Description of Artificial Sequence:
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                                    TYPE: RNA
ORGANISM: Artificial Sequence
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Best Local Similarity 100.0
Matches 19; Conservative
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APPLICANT: Addnav, Vasant
APPLICANT: Vaish, Naredara
APPLICANT: Weswiggen, James
TITLE OF INVENTION: RNA Interference Mediated Inhibition Of Vascular Endothelial
TITLE OF INVENTION: Gene Expression Using Short Interfering Nucleic Acid (sinA)
TITLE OF INVENTION: Gene Expression Using Short Interfering Nucleic Acid (sinA)
TITLE OF INVENTION: Gene Expression Using Short Interfering Nucleic Acid (sinA)
TITLE OF INVENTION WIMBER: US 10/844,016
CURRENT FILING DATE: 2004-09-16
PRIOR FILING DATE: 2004-04-23
PRIOR FILING DATE: 2004-04-23
PRIOR PLILING DATE: 2004-01-26
PRIOR APPLICATION NUMBER: US 10/66,255
PRIOR FILING DATE: 2003-09-16
PRIOR FILING DATE: 2003-09-16
PRIOR PLILING DATE: 2003-09-16
PRIOR PLILING DATE: 2003-09-16
PRIOR APPLICATION NUMBER: US 10/66,255
PRIOR APPLICATION NUMBER: US 10/66,255
PRIOR APPLICATION NUMBER: US 00/330/0502
PRIOR PRILING DATE: 2003-07-03
PRIOR PLILING DATE: 2003-07-03
PRIOR PLILING DATE: 2003-07-03
PRIOR PLILING DATE: 2003-07-03
PRIOR PRILING DATE: 2003-07-03
PRIOR PLILING DATE: 2003-07-03
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                         PRIOR APPLICATION NUMBER: US 60/399,348
PRIOR FILING DATE: 2002-07-29
PRIOR FILING DATE: 2002-07-29
PRIOR PILING DATE: 2004-05-24
Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 4252
SOFTWARE: Patentin version 3.3
SEQ ID NO 2809
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OTHER INFORMATION: Phosphorothioate 3'-Internucleotide Linkage
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Pred. No. 8.4;
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CTHER INFORMATION: X stands for nitroindole universal base US-10-944-611-2809
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ORGANISM: Artificial Sequence
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FILING DATE: 2002-07-03
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NAME/KEY: misc_feature
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Best Local Similarity
Matches 15; Conserva
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Gaps . ,

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APPLICANT: McSwiggen, James TITLE OF INVENTION: NA Interference Mediated Inhibition Of Vascular Endothelial TITLE OF INVENTION: NA Interference Mediate Endothelial Growth Factor Recept TITLE OF INVENTION: Growth Factor And Vascular Endothelial Growth Factor Recept TITLE OF INVENTION: Gene Expression Using Short Interfering Nucleic Acid (siNA) PITLE REFERENCE: 400/235 (MRHBO2-742-S)
CURRENT APPLICATION NUMBER: US/10/944,611
CURRENT FILING DATE: 2004-09-16
                                                                   PRIOR APPLICATION NUMBER: US 10/844,076
PRIOR PAPLICATION NUMBER: US 10/844,076
PRIOR PELICATION NUMBER: US 10/831,620
PRIOR PELICATION NUMBER: US 10/831,620
PRIOR PILING DATE: 2004-04-23
PRIOR PILING DATE: 2003-09-23
PRIOR PELICATION NUMBER: US 10/665,255
PRIOR PILING DATE: 2003-09-16
PRIOR PILING DATE: 2002-07-03
PRIOR PILING DATE: 2002-07-03
PRIOR PILING DATE: 2002-07-29
PRIOR PILING DATE: 2002-07-29
PRIOR PILING DATE: 2004-05-24
PRIOR PILING DATE: 2004-05-34
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LOCATION: (1)._(1)
OTHER INFORMATION: 5'-3 attached terminal deoxyabasic moeity
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PRIOR APPLICATION NUMBER: US 10/844,076
PRIOR FILLING DATE: 2004-05-11
PRIOR PLICATION NUMBER: US 10/831,620
PRIOR PLICATION NUMBER: US 10/764,957
PRIOR PLICATION NUMBER: US 10/764,957
PRIOR PLICATION NUMBER: US 10/670,011
PRIOR PLICATION NUMBER: US 10/670,011
PRIOR PLICATION NUMBER: US 10/670,011
PRIOR PLICATION NUMBER: US 10/65,255
PRIOR FILLING DATE: 2003-09-16
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PRIOR APPLICATION NUMBER: US 10/664,767
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 CUGAGUUUAAAAGGCACCC 19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: Artificial Sequence
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Best Local Similarity 78.9'
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-10-944-611-3141
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: RNA
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APPLICANT: Vaich, Narendra
APPLICANT: Vaich, Narendra
APPLICANT: Walch, Narendra
TTTLE OF INVENTION: Growth Factor And Vascular Endothelial
TTTLE OF INVENTION: Growth Factor And Vascular Endothelial Growth Factor Receptor
TTTLE OF INVENTION: Growth Factor And Vascular Endothelial Growth Factor Receptor
TTTLE OF INVENTION: Growth Factor And Vascular Endothelial Growth Factor Receptor
TTTLE OF INVENTION: Growth Factor And Vascular Endothelial Growth Factor Receptor
TTTLE OF INVENTION: Growth Factor And Vascular Endothelial Growth Factor Receptor
TTTLE OF INVENTION WUMBER: US 10/944,611
CURRENT FILING DATE: 2004-09-16
PRIOR PILING DATE: 2004-01-26
PRIOR FILING DATE: 2004-01-26
PRIOR FILING DATE: 2004-01-26
PRIOR FILING DATE: 2004-01-26
PRIOR FILING DATE: 2003-09-16
PRIOR PLICATION WUMBER: US 10/66,255
PRIOR APPLICATION WUMBER: US 10/66,767
PRIOR FILING DATE: 2003-09-16
PRIOR FILING DATE: 2003-09-16
PRIOR FILING DATE: 2003-09-16
PRIOR FILING DATE: 2003-09-16
PRIOR FILING DATE: 2003-09-23
PRIOR PLICATION WUMBER: US 60/393,796
PRIOR FILING DATE: 2003-09-23
PRIOR APPLICATION WUMBER: US 60/393,48
PRIOR APPLICATION WUMBER: US 60/393,48
PRIOR APPLICATION WUMBER: US 60/393,48
PRIOR PRILING DATE: 2004-02-29
PRIOR FILING DATE: 2004-02-29
PRIOR FILING DATE: 2004-02-29
PRIOR FILING DATE: 2004-02-39
PRIOR FILING DATE: 2004-05-24
PRIOR FILING DATE: 2004-05-34
PRIOR 
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Gene Expression Using Short Interfering Nucleic Acid (siNA)
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APPLICANT: Vaish, Nacendra
APPLICANT: Vaish, Nacendra
APPLICANT: Vaish, Nacendra
APPLICANT: Wish, Nacendra
APPLICANT: McSwiggen, James
TITLE OF INVENTION: Growth Factor And Vascular Endochelial Growth Factor Receptor
TITLE OF INVENTION: Gene Expression Using Short Interfering Nucleic Acid (siNA)
FILE REPERENCE: 400/235 (MBHB02-742-S)
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100.0%; Pred. No. 8.4;
tive 0; Mismatches 0.
                                                                                                                                                                                         Sequence 3139, Application US/10944611
GENERAL INFORMATION:
APPLICANT: Sirna Therapeutics, Inc.
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   CTGAGTTTAAAAGGCACCC 1
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Kossen, Karl
Zinnen, Shawn
Vaish, Narendra
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APPLICANT: Kossen, Karl
APPLICANT: Zinnen, Shawn
APPLICANT: Vaish, Narendra
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Best Local Similarity 100.
Matches 19; Conservative
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US-10-944-611-3141/C
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LENGTH: 19
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siNA sense region

Gaps

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APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Richards, Ivan
APPLICANT: Richards, Ivan
APPLICANT: McSwiggen, James
TITLE OF INVENTION: RNA Interference Mediated Inhibition Of Vascular Endothelial
TITLE OF INVENTION: Growth Factor And Vascular Endothelial Growth Factor Receptor
TITLE OF INVENTION: Gene Expression Using Short Interfering Nucleic Acid (siNA)
FILE REPERENCE: 400/236 (MBHB02-742-U)
CURRENT APPLICATION NUMBER: US 10/962,898
CURRENT FILING DATE: 2004-10-12
PRIOR PILING DATE: 2004-09-16
PRIOR FILING DATE: 2004-09-16
PRIOR FILING DATE: 2004-05-11
                                                                                                                                                                                                                                                                                                                                                                                                                                  OTHER INFORMATION: Description of Artificial Sequence: sinh sense region
              PRIOR FILING DATE: 2003-02-20
PRIOR PILING DATE: 2003-02-20
PRIOR APPLICATION NUMBER: US 60/399,796
PRIOR FILING DATE: 2002-07-03
PRIOR FILING DATE: 2002-07-29
PRIOR PILING DATE: 2002-07-29
PRIOR PILING DATE: 2004-05-24
Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 4252
SOFTWARE: PatentIn version 3.3
SEQ ID NO 3144
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APPLICATION NUMBER: PCT/US03/05022
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OTHER INFORMATION: 2'-deoxy-2'-fluoro
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OTHER INFORMATION: 2'-deoxy-2'-fluoro
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NAME/KEY: misc_feature
NOCATION: (18)..(19)
OTHER INFORMATION: 2'-O-methyl
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OTHER INFORMATION: 2'-0-methyl
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LOCATION: (5)...(5)
OTHER INFORMATION: 2'-O-methyl
FEATURE:
NAME/KEY: misc_feature
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LOCATION: (12)...(14)
OTHER INFORMATION: 2'-0-methyl
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Best Local Similarity 78.33
Best Local Similarity 78.33
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NAME/KEY: misc_feature
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APPLICANT: Jadhav, Vasant
APPLICANT: Jadhav, Vasant
APPLICANT: Jadhav, Vasant
APPLICANT: Jadhav, Vasant
APPLICANT: Stanen, Shawn
APPLICANT: Zinnen, Shawn
APPLICANT: Vaish, Narendra
APPLICANT: Waish, Narendra
TITLE OF INVENTION: Growth Factor And Vascular Endothelial
TITLE OF INVENTION: ADMER: US/10/944,611
CURRENT APPLICATION NUMBER: US 10/844,016
FRIOR FILING DATE: 2004-09-16
FRIOR FILING DATE: 2004-04-23
FRIOR FILING DATE: 2003-09-16
FRIOR FILING DATE: 2003-09-18
FRIOR FILING DATE: 2003-09-18
FRIOR FILING DATE: 2003-09-16
FRIOR FILING DATE: 2003-09-16
FRIOR FILING DATE: 2003-09-16
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                PRIOR PAPLICATION NUMBER: PCT/USO3/05022
PRIOR PILING DATE: 2003-02-20
PRIOR PLILING DATE: 2003-07-03
PRIOR PLLING DATE: 2002-07-03
PRIOR PLLING DATE: 2002-07-29
PRIOR PLLING DATE: 2002-07-29
PRIOR APPLICATION NUMBER: PCT/USO4/16390
PRIOR PLILING DATE: 2004-05-24
Remaining Prior Application data removed - 8
NUMBER OF SEQ ID NOS: 4252
SOFTWARE: PatentIn version 3.3
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NAME/KEY: misc_feature
LOCATION: (17)..(19)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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LOCATION: (1). (2)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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LOCATION: (6)._(8)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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OTHER INFORMATION: 2'-deoxy-2'-fluoro
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Sequence 3144, Application US/10944611
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                  TYPE: RNA
ORGANISM: Artificial Sequence
FILING DATE: 2003-09-16
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Matches 19; Conservative
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NAME/KEY: misc_feature
LOCATION: (15)..(15)
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APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Richards, Ivan
APPLICANT: Richards, Ivan
APPLICANT: McSwidgen, James
TITLE OF INVENTION: RNA Interference Mediated Inhibition Of Vascular Endothelial
TITLE OF INVENTION: Growth Factor And Vascular Endothelial Growth Factor Recept
TITLE OF INVENTION: Gene Expression Using Short Interfering Nucleic Acid (siNA)
FILE REPERENCE: 400/236 (MBHB02-742-U)
CURRENT APPLICATION NUMBER: 2004-10-12
                                                                                                                                                                                                                                                                                                                                                                                                                              OTHER INFORMATION: Description of Artificial Sequence: sinA sense region
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PRIOR PILING DATE: 2004-09-16

PRIOR PILING DATE: 2004-09-16

PRIOR PILING DATE: 2004-09-16

PRIOR PILING DATE: 2004-06-13

PRIOR APPLICATION NUMBER: US 10/841,620

PRIOR PILING DATE: 2004-04-23

PRIOR PILING DATE: 2004-01-26

PRIOR PILING DATE: 2003-09-16

PRIOR PILING DATE: 2002-07-03

PRIOR PILING DATE: 2002-07-03
PRIOR APPLICATION NUMBER: US 60/393,796
PRIOR FILING DATE: 2002-07-03
PRIOR APPLICATION NUMBER: US 60/399,348
PRIOR FILING DATE: 2002-07-29
Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 4252
SOFTWARE: Patentin version 3.3
SEQ ID NO 2967
LENGTH: 19
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: LOCATION: (1)...(1)

: OTHER INPORMATION: 5'-3 attached terminal deoxyabasic moeity

US-10-962-898-2967
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8.4;
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Matches 19; Conservative
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                                                                                                                                                                                                                                                                                                                           TYPE: RNA
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                                                                                                                                                                                                                                                                                                                                                                                              FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FEATURE:
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APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Richards, Ivan
APPLICANT: Richards, Ivan
TITLE OF INVENTION: Rowth Factor And Vascular Endothelial
TITLE OF INVENTION: Growth Factor And Vascular Endothelial Growth Factor Receptor
TITLE OF INVENTION: Gene Expression Using Short Interfering Nucleic Acid (siNA)
FILE REPRENCE: 400/236 (MBHB02-742-U)
CURRENT APPLICATION NUMBER: US/10/962,898
CURRENT APPLICATION NUMBER: US 10/944,644
PRIOR FILING DATE: 2004-09-16
PRIOR FILING DATE: 2004-05-11
PRIOR FILING DATE: 2004-05-11
PRIOR FILING DATE: 2004-04-23
PRIOR FILING DATE: 2004-04-23
PRIOR FILING DATE: 2004-01-26
PRIOR FILING DATE: 2004-09-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OTHER INFORMATION: Description of Artificial Sequence: sina antisense region
                                                                      PRIOR APPLICATION NUMBER: US 10/764,957
PRIOR APPLICATION NUMBER: US 10/670,011
PRIOR FILING DATE: 2003-09-23
PRIOR FILING DATE: 2003-09-23
PRIOR PILING DATE: 2003-09-16
PRIOR PILING DATE: 2003-02-07
PRIOR PILING DATE: 2003-02-07
PRIOR PILING DATE: 2002-07-03
PRIOR PILING DATE: 2002-07-03
PRIOR PILING DATE: 2002-07-29
PRIOR PILING DATE: 2002-07-29
PRIOR PILING DATE: 2002-07-29
PRIOR PILING DATE: 2002-07-29
PRIOR PILING DATE: 2003-03-33
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LOCATION: (20)...(20)
OTHER INFORMATION: Phosphorothioate 3'-Internucleotide Linkage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 14; Length 19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME/KEY: misc_feature
i LOCATION: (20)..(20)
i CTHER INFORMATION: X stands for nitroindole universal base
US-10-962-898-2809
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                                          R FILING DATE: 2004-04-23

R APPLICATION NUMBER: US 10/764,957

R APPLICATION NUMBER: US 10/670,011

R APPLICATION NUMBER: US 10/665,255

R FILING DATE: 2003-09-23
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Pred. No.
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APPLICATION NUMBER: PCT/US03/05022
FILING DATE: 2003-02-20
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APPLICATION NUMBER: US 10/664,767
       APPLICATION NUMBER: US 10/831,620
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ORGANISM: Artificial Sequence
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Best Local Similarity
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LENGTH: 19
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OTHER INFORMATION: Description of Artificial Sequence: sinA antisense region
                                                                                 DB 14; Length 19;
                                                                              90.5%; Score 19;
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US-10-962-898-3141
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
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GENERAL INFORMATION:

GENERAL INFORMATION:

APPLICANT: Sidnar Therapeutics, Inc.

APPLICANT: Sidnards, Ivan

APPLICANT: Richards, Ivan

APPLICANT: Michards, Ivan

APPLICANT: Michards, Ivan

ITILE OF INVENTION: Mames

ITILE OF INVENTION: Growth Factor And Vascular Endothelial

ITILE OF INVENTION NUMBER: US 10/944,644

FRIOR APPLICATION NUMBER: US 10/644,076

FRIOR APPLICATION NUMBER: US 10/664,767

FRIOR APPLICATION NUMBER: US 10/664,767

FRIOR APPLICATION NUMBER: US 60/399,348

FRIOR APPLICATION NUMBER: US 60/399,348
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GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Richards, Ivan
TITLE OF INVENTION: RNA Interference Mediated Inhibition Of Vascular Endothelial
TITLE OF INVENTION: Growth Factor And Vascular Endothelial Growth Factor Receptor
TITLE OF INVENTION: Gene Expression Using Short Interfering Nucleic Acid (sinA)
FILE REFERENCE: 400/236 (MBHB02-742-U)
CURRENT APPLICATION NUMBER: US/10/962,898
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                                 Gaps
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                                 0; Indels
                              Mismatches
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NUMBER OF SEQ ID NOS: 4252
SOFTWARE: PatentIn version 3.3
SEQ ID NO 3139
LENGTH: 19
   78.9%; Pred. No.
                                 4;
                                                                                       1 CUGAGUUUAAAAGGCACCC 19
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                                                                                                                     TYPE: RNA
ORGANISM: Artificial Sequence
Best Local Similarity 78.9
Matches 15; Conservative
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hes 19; Conserva
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US-LU-98c-898-914, Application US/10962898

Sequence 3143, Application US/10962898

SERVERAL INFORMATION:
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Richards, Ivan
TITLE OF INVENTION: RNA Interference Mediated Inhibition Of Vascular Endothelial
TITLE OF INVENTION: Growth Factor And Vascular Endothelial Growth Factor Receptor
TITLE OF INVENTION: Growth Factor And Vascular Endothelial Growth Factor Receptor
TITLE OF INVENTION: Growth Factor And Vascular Endothelial Growth Factor Receptor
TITLE OF INVENTION: Growth Factor And Vascular Endothelial Growth Factor Roughest Application NUMBER: US/10/962,898
CURRENT APPLICATION NUMBER: US 10/944,644
FRIOR APPLICATION NUMBER: US 10/944,644
FRIOR APPLICATION NUMBER: US 10/944,076
FRIOR APPLICATION NUMBER: US 10/944,957
FRIOR APPLICATION NUMBER: US 10/64,957
FRIOR APPLICATION NUMBER: US 10/64,957
FRIOR PILING DATE: 2004-01-26
FRIOR FILING DATE: 2003-09-23
FRIOR FILING DATE: 2003-09-23
FRIOR FILING DATE: 2003-09-16
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FRIOR APPLICATION NUMBER: US 10/844,076

PRIOR APPLICATION NUMBER: US 10/831,620

PRIOR FILING DATE: 2004-06-13

PRIOR FILING DATE: 2004-06-13

PRIOR PELING DATE: 2004-01-26

PRIOR PELING DATE: 2004-01-26

PRIOR PELING DATE: 2004-01-26

PRIOR PELING DATE: 2003-09-23

PRIOR PELING DATE: 2003-09-16

PRIOR PELING DATE: 2002-07-03

PRIOR PELING DATE: PAPALICATION NUMBER: US 60/399,348

PRIOR PELING DATE: 2002-07-03

PRIOR PELING DATE: PAPALICATION NUMBER: US 60/399,348

PRIOR PELING DATE: 2002-07-03

PRIOR PELING DATE: PAPALICATION NUMBER: US 60/399,348

PRIOR PELING DATE: PAPALICATION NUMBER: US 60/399,348
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OTHER INFORMATION: 5'-3 attached terminal deoxyabasic moeity
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Matches 15; Conservative
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APPLICANT: Jadhav, Vasant
APPLICANT: Jadhav, Vasant
APPLICANT: Zinner, Sham
APPLICANT: Zinner, Sham
APPLICANT: Zinner, Sham
APPLICANT: Vaish, Narendra
APPLICANT: Waish, Narendra
APPLICANT: Waish, Narendra
APPLICANT: McSwiggen, James
TITLE OF INVENTION: Gene Expression Using Short Interfering Nucleic Acid (sina)
FILE REPERENCE: 400/235 (MBH802-742-8)
CURRENT APPLICATION NUMBER: US/10/944,611
CURRENT FILING DATE: 2004-09-16
PRIOR APPLICATION NUMBER: US 10/844,076
PRIOR PLILNG DATE: 2004-04-23
PRIOR FILING DATE: 2004-04-23
                                                                                                                                                                                                                                                 OTHER INFORMATION: Description of Artificial Sequence: sinA sense region
  PRIOR FILING DATE: 2002-07-29
Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 4252
SOCTWARE: PatentIn version 3.3
SEQ ID NO 3144
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APPLICATION NUMBER: US 10/764,957
FILING DATE: 2004-01-26
APPLICATION NUMBER: US 10/670,011
FILING DATE: 2003-09-23
                                                                                                                                                                                                                                                                                                                                                                                                             NAME/KEY: misc_feature
LOCATION: (4)._(4)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME/KEY: misc feature
LOCATION: (6). (11)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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OTHER INFORMATION: 2'-deoxy-2'-fluoro
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OTHER INFORMATION: 2'-0-methyl
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OTHER INFORMATION: 2'-0-methyl
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OTHER INFORMATION: 2'-0-methyl
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LOCATION: (18)..(19)
CHER INFORMATION: 2'-O-methyl
US-10-962-898-3144
                                                                                                                                                                                              ORGANISM: Artificial Sequence
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Best Local Similarity 78.9°
Matches 15; Conservative
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                                                                                                                                                                          TYPE: RNA
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APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Richards, Ivan
APPLICANT: Richards, Ivan
APPLICANT: McSwiggen, James
TITLE OF INVENTION: RNA Interference Mediated Inhibition Of Vascular Endothelial
TITLE OF INVENTION: Growth Factor And Vascular Endothelial Growth Factor Receptor
TITLE OF INVENTION: Growth Factor Using Short Interfering Nucleic Acid (siNA)
TITLE OF INVENTION: Growth Factor May Secular Endothelial Growth Factor Receptor
TITLE OF INVENTION: Growth Factor And Vascular Endothelial
TITLE OF INVENTION: Growth Factor And Vascular Endothelial
CURRENT FILING DATE: 2004-01-12
PRIOR APPLICATION NUMBER: US 10/844,076
PRIOR FILING DATE: 2004-04-23
PRIOR FILING DATE: 2004-04-23
PRIOR FILING DATE: 2003-09-16
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                      PRIOR APPLICATION NUMBER: US 60/393,796
PRIOR FILING DATE: 2002-07-03
PRIOR APPLICATION NUMBER: US 60/399,348
PRIOR PILING DATE: 2002-07-29
Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 4252
SOFTWARE: Patentin version 3.3
SEQ ID NO 3143
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ö
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APPLICATION NUMBER: PCT/US03/05022
FILING DATE: 2003-02-20
APPLICATION NUMBER: US 60/393,796
FILING DATE: 2002-07-03
APPLICATION NUMBER: US 60/399,348
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; LOCATION: (17)...(19)
; OTHER INFORMATION: 2'-deoxy-2'-fluoro
US-10-962-898-3143
                                                                                                                                                                                                                                                                                                                                                                                                          NAME/KEY: misc_feature
LOCATION: (1)._(2)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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OTHER INFORMATION: 2'-deoxy-2'-fluoro
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LOCATION: (15) ... (15)
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                                                                                                                                                                                                                                                                          TYPE: RNA
ORGANISM: Artificial Sequence
FILING DATE: 2003-02-20
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Best Local Similarity 100.
Matches 19; Conservative
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NAME/KEY: misc_feature
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US-10-962-898-3144/c
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PRIOR APPLICATION NUMBER: US 10/665,255
PRIOR APPLICATION NUMBER: US 10664,767
PRIOR APPLICATION NUMBER: US 101664,767
PRIOR APPLICATION NUMBER: PCT/US03/05022
PRIOR PILING DATE: 2003-02-20
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PRIOR APPLICATION NUMBER: US 60/399,348
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PRIOR APPLICATION POS: 4222
SOCTWARE: PRIOR PRIO
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Search completed: August 19, 2005, 00:02:13 Job time : 1736 secs

Dp

Pending Nucleic Acid and Pending Amino Acid database searches generate two sets of results each. The Pending databases have been split into two parts to reduce the amount of time required for their daily updates. This results in more machine time being available for processing searches.

Searches run against the Nucleic Acid Pending database produce two sets of results, with the extensions .rnpm and .rnpn

Searches run against the Amino Acid Pending database produce two sets of results, with the extensions .rapm and .rapn

Because they contain data that is confidential, the results of Pending database searches should not be left in the case.

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us-10-665-951-2185.rnpm

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GenCore version 5.1.6
Copyright (c) 1993 - 2005 Compugen Ltd.
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq length: 30
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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	qı	8 US-10-664-668-231	8 US-10-665-951-	2 US-10-758-155-231	3 US-10-831-620-231	3 US-10-844-076-231	PCT-US03-05028-39	PCT-US03-05028-39	PCT-US03-05028-39	PCT-US03-05028-40	PCT-US03-05028-44	PCT-US03-05028-45	PCT-US03-05028-4	PCT-US03-05028-45	PCT-US03-05028-45	PCT-US03-05028-46	PCT-US03-05028-48	PCT-US03-05028-48	PCT-US03-05346-29	PCT-US03-05346-30	PCT-US03-05346-30	PCT-US03-05346-34	4 US-10-444-853A-2	4 US-10-444-853A-3	4 US-10-444-853A-3	4 US-10-444-853A-3	4 US-10-444-853A-6	4 US-10-444-853A-6	8 US-10-664-668-20	8 US-10-664-668-2	8 US-10-664-668-21	8 US-10-664-668-216	8 US-10-664-668-21	8 US-10-664-668-218	8 US-10-664-668-221	8 US-10-664-668-222	8 US-10-664-668-22	8 US-10-664-668-22	8 US-10-664-668-2	8 US-10-664-668-22						
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ALIGNMENTS

APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: McSwiggen, James
APPLICANT: McSwiggen, James
APPLICANT: Beigelman, Leonid
APPLICANT: Pavco, Pamela
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Vascular Endothelial
TITLE OF INVENTION: Geowth Factor and Vascular Endothelial Growth Factor Receptor
TITLE OF INVENTION: Gene Expression Using Short Interfering Nucleic Acid (siNA) OTHER INFORMATION: Description of Artificial Sequence: sinA antisense region ö Gaps ö NAME/KEY: misc feature LOCATION: (20) ... (20) OTHER INFORMATION: Phosphorothioate 3'-Internucleotide Linkage 90.5%; Score 19; DB 58; Length 19; 78.9%; Pred. No. 12; tive 4; Mismatches 0; Indels LOCATION: (20) ... (20) ... (20) ... OTHER INFORMATION: X stands for nitroindole universal base US-10-664-668-2313 Sequence 2313, Application US/10664668; GENERAL INFORMATION: ; Sequence 2313, Application US/10665951; GENERAL INFORMATION: 1 CUGAGUUUAAAAGGCACCC 19 TYPE: RNA ORGANISM: Artificial Sequence 15; Conservative FEATURE:
NAME/KEY: misc_feature Query Match Best Local Similarity US-10-664-668-2313/c US-10-665-951-2313/c 19 Matches g ð

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APPLICANT: Beigelman, Leonid
APPLICANT: Beigelman, Leonid
APPLICANT: Beigelman, Leonid
APPLICANT: Beigelman, Leonid
APPLICANT: Beyoo, Pameala
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Vascular Endothelial
TITLE OF INVENTION: Gene Expression Using Short Interfering Nucleic Acid (siNA)
TITLE OF INVENTION: Gene Expression Using Short Interfering Nucleic Acid (siNA)
TITLE OF INVENTION: Gene Expression Using Short Interfering Nucleic Acid (siNA)
TITLE OF INVENTION: Government Us/10/831,620
CURRENT FILING DATE: 2004-04-23
PRIOR APPLICATION NUMBER: US 10/764,957
PRIOR APPLICATION NUMBER: US 10/759,803
PRIOR PILING DATE: 2004-01-12
PRIOR PELING DATE: 2004-01-12
PRIOR PELING DATE: 2003-11-24
PRIOR FILING DATE: 2003-11-24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OTHER INFORMATION: Description of Artificial Sequence: sinA antisense region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Remaining Prior Application data removed - See File Wrapper or PALM. NUMBER OF SEQ ID NOS: 2751
SOFTWARE: PatentIn version 3.3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME/KEY: misc feature LOCATION: (20) ... (20) OTHER INFORMATION: Phosphorothioate 3'-Internucleotide Linkage
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; LOCATION: (20)..(20)
; OTHER INFORMATION: X stands for nitroindole universal base
US-10-758-155-2313
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Pred. No. 12;
4; Mismatches
                                  PRIOR FILING DATE: 2002-07-03
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PRIOR FILING DATE: 2002-11-04
PRIOR APPLICATION NUMBER: US 10/306,747
PRIOR FILING DATE: 2002-11-27
PRIOR APPLICATION NUMBER: PCT/US 02/17674
PRIOR PLING DATE: 2002-05-29
PRIOR PLING DATE: 2002-05-29
PRIOR PLING DATE: 2002-02-20
PRIOR APPLICATION NUMBER: US 60/363,124
PRIOR PLING DATE: 2002-02-20
            APPLICATION NUMBER: US 60/393,796
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ORGANISM: Artificial Sequence
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Best Local Similarity 78.9%;
Matches 15; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-10-831-620-2313/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: Pater
SEQ ID NO 2313
LENGTH: 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FEATURE:
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TITLE OF INVENTION: RNA Interference Mediated Inhibition of Vascular Endothelial
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Vascular Endothelial
TITLE OF INVENTION: Growth Factor and Vascular Endothelial Growth Factor Receptor
TITLE OF INVENTION: Gene Expression Using Short Interfering Nucleic Acid (sinh)
FILE REFERENCE: 400/141 (MEHB02742-N)
CURRENT APPLICATION NUMBER: US/07/58,155
PRIOR PALICATION NUMBER: US/0665,951
PRIOR PALICATION NUMBER: US/0665,951
PRIOR PALICATION NUMBER: US/0664,668
PRIOR PILING DATE: 2003-09-18
PRIOR APPLICATION NUMBER: PCT/US 03/05022
PRIOR PILING DATE: 2003-09-20
PRIOR FILING DATE: 2003-02-20
PRIOR FILING DATE: 2003-02-20
PRIOR FILING DATE: 2003-02-20
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                 CURRENT APPLICATION NUMBER: US 10/65,951

CURRENT APPLICATION NUMBER: US 10/664,668

PRIOR PLILING DATE: 2003-09-18

PRIOR PLILING DATE: 2003-09-18

PRIOR PLILING DATE: 2003-09-28

PRIOR PLILING DATE: 2003-09-28

PRIOR PLILING DATE: 2003-02-20

PRIOR PLING DATE: 2002-07-29

PRIOR PLILING DATE: 2002-11-04

PRIOR PLILING DATE: 2002-11-24

PRIOR PLILING DATE: 2002-11-27

PRIOR PLILING DATE: 2002-11-27

PRIOR PLILING DATE: 2002-11-27

PRIOR PLILING DATE: 2002-11-27

PRIOR PLILING DATE: 2002-05-29

PRIOR PLILING DATE: 2002-05-39

PRIOR PLILING DATE: 2002-05-06-06

PRIOR PLILING DATE: 2002-06-06

PRIOR PLILING DATE: 2002-06-06

PRIOR PLILING DATE: 2002-06-06

PRIOR PLING DATE: 2002-06-06

PRIOR PLILING DATE: 2002-06-06

PRIOR PRIOR PLILING DATE: 2002-06-06

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OTHER INFORMATION: Phosphorothioate 3'.Internucleotide Linkage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 19;
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; LOCATION: (20)...(20)

; OTHER INFORMATION: X stands for nitroindole universal base

US-10-665-951-2313
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(MBHB02-742-F)
3ER: US/10/665,951
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       19 CTGAGTTTAAAAGGCACCC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: RNA
ORGANISM: Artificial Sequence
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Best Local Similarity
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Gaps

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Indels

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OTHER INFORMATION: Description of Artificial Sequence: sinA sense region
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; OTHER INFORMATION: Phosphorothioate 3'-Internucleotide Linkage
PCT-US03-05028-390
            LOCATION: (20)...(20)
OTHER INFORWATION: Phosphorothioate 3'-Internucleotide Linkage PEATURE:
NAME/KEY: misc_feature
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME/KEY: misc feature
LOCATION: (1). [5)
OTHER INFORMATION: Phosphorothioate 3'-Internucleotide Linkage
                                                                                                                    63; Length 19;
                                                       LOCATION: (20)...(20)

COTHER INFORMATION: X stands for nitroindole universal base US-10-844-076-2313
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                                                                                                                                              Indels
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                                                                                                                   Query Match 90.5%; Score 19; DB Best Local Similarity 78.9%; Pred. No. 12; Matches 15; Conservative 4; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       OTHER INFORMATION: n stands for thymidine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    90.5%; Score 19;
                                                                                                                                                                                                                                                                  Sequence 390, Application PC/TUS0305028 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                            APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: McSwiggen, James
APPLICANT: Chowilta
APPLICANT: Chowilta, Bharat
                                                                                                                                                                         1 CUGAGUTUAAAAGGCACCC 19
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NAME/KEY: misc_feature
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SOFTWARE: Pate
SEQ ID NO 390
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OTHER INFORMATION: Description of Artificial Sequence: sinA antisense region
FEATURE:
                                                                                                                    OTHER INFORMATION: Description of Artificial Sequence: sinA antisense region
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     Remaining Prior Application data removed - See File Wrapper or PALM. NUMBER OF SEQ ID NOS: 2751
SOFTWARE: Patentin version 3.3
SEQ ID NO 2313
LENGTH: 19
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                                                                                                                                 FEATURE:
NAME/KEY: misc_feature
LOCATION: (20)...(20)
OTHER INFORWATION: Phosphorothioate 3'-Internucleotide Linkage
                                                                                                                                                                                                                                                                  Query Match 90.5%; Score 19; DB 63; Length 19; Best Local Similarity 78.9%; Pred. No. 12; Matches 15; Conservative 4; Mismatches 0; Indels
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NAME/KEY: misc_feature

LOCATION: (20)...(20)

OTHER INFORMATION: X stands for nitroindole universal base
US-10-831-620-2313
                                                                                                                                                                                                                                                                                                                        1 CUGAGUUUAAAAGGCACCC 19
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                                                                            TYPE: RNA
ORGANISM: Artificial Sequence
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ORGANISM: Artificial Sequence
PRIOR FILING DATE: 2003-09-16
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Sequence 398, Application PC/TUS0305028 GENERAL INFORMATION:
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Jamison, Sharon
Usman, Nassim
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TITLE OF INVENTION: RNA Interference Mediated Inhibition of Gene Expréssion Using
TITLE OF INVENTION: Short Interfering Nucleic Acid (sina)
FILE REFERENCE: 400/104 (MBBB 03-198)
CURRENT APPLICATION NUMBER: PCT/US03/05028
CURRENT FILING DATE: 2003-02-20
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LOCATION: (20) . (20)
OTHER INFORMATION: Phosphorothioate 3'-Internucleotide Linkage
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OTHER INFORMATION: Phosphorothioate 3'-Internucleotide Linkage
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                           Indels
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100.0%; Pred. No. 13;
tive 0; Mismatches
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PRIOR APPLICATION NUMBER: US 60/363,124
PRIOR FILING DATE: 2002-03-11
PRIOR PELING DATE: 2002-03-11
PRIOR PELING DATE: 2002-02-05
PRIOR PILING DATE: 2002-06-06
PRIOR PILING DATE: 2002-06-06
PRIOR PILING DATE: 2002-06-06
PRIOR FILING DATE: 2002-06-06
PRIOR FILING DATE: 2002-08-29
PRIOR PILING DATE: 2002-09-05
PRIOR PELING DATE: 2002-09-05
PRIOR PELING DATE: 2002-09-05
PRIOR PELING DATE: 2002-09-09
PRIOR PELING DATE: 2002-09-09
PRIOR PILING DATE: 2002-09-09
PRIOR PILING DATE: 2002-09-09
PRIOR PILING DATE: 2002-09-09
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NAME/KEY: misc_feature
LOCATION: (20)..(21)
OTHER INFORMATION: n stands for thymidine
                                                                                                                                                                                                              PCT-US03-05028-394/c; Sequence 394, Application PC/TUS0305028; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                    APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: McSwiggen, James
APPLICANT: Beigelman, Leonid
APPLICANT: Chowrira, Bharat
                                                                                                     CUGAGUUUAAAAGGCACCC 19
                                                                   1 CUGAGUUDAAAAGGCACCC 19
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SOFTWARE: Patentin version 3.2
SEQ ID NO 394
LENGTH: 21
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Fosnaugh, Kathy
Jamison, Sharon
Best Local Similarity 100.
Matches 19; Conservative
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Matches 15, Conservative
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APPLICANT: Thompson, James
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Gene Expression Using
TITLE OF INVENTION: Short Interfering Nucleic Acid (sina)
FILE REFERENCE: 400/104 (MBHB 03-198)
CURRENT APPLICATION NUMBER: PCT/US03/05028
CURRENT FILING DATE: 2003-02-20
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LOCATION: (1). \( (4)\) OTHER INFORMATION: Phosphorothioate 3'-Internucleotide Linkage
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OTHER INFORMATION: Phosphorothioate 3'-Internucleotide Linkage
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13;
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                                                                                               ; LOCATION: (20) ... (21)
PO THER INFORMATION: n stands for thymidine PCT-US03-05028-398
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FEATURE:
NAME/KEY: misc_feature
LOCATION: (15)..(15)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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LOCATION: (1). (2)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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COCATION: (6). (8)
THER INFORMATION: 2'-deoxy-2'-fluoro
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ORGANISM: Artificial Sequence
FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME/KEY: misc_feature
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Best Local Similarity
Matches 19; Conserv
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1 CUGAGUUUAAAAGGCACCC 19

RESULT 8 PCT-US03-05028-398

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TITLE OF INVENTION: RNA Interference Mediated Inhibition of Gene Expression Using TITLE OF INVENTION: Short Interfering Nucleic Acid (siNA)

TITLE OF INVENTION: Short Interfering Nucleic Acid (siNA)

FILE REFERENCE: 400/104 (MBHB 23.198)

CURRENT APPLICATION NUMBER: DC 50/363,124

FRIOR PELING DATE: 2002-02-01

PRIOR PELING DATE: 2002-02-01

PRIOR PELING DATE: 2002-02-05

PRIOR APPLICATION NUMBER: US 60/386,782

PRIOR APPLICATION NUMBER: US 60/406,784

PRIOR FILING DATE: 2002-08-05

PRIOR FILING DATE: 2002-09-09

PRIOR FILING DATE: 2002-09-09

PRIOR PELING DATE: 2002-09-09

PRIOR PELING DATE: 2002-09-09

PRIOR PELING DATE: 2003-09-09

PRIOR PELING DATE: 2003-09-15

PRIOR FILING DATE: 2003-09-15

NUMBER OF SEQ ID NOS: 932

SOFTWARE: Patentin version 3.2
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TITLE OF INVENTION: Short interference Mediated Inhibition of Gene Expression Using
TITLE OF INVENTION: Short interfering Nucleic Acid (sinA)
TITLE OF INVENTION: Short interfering Number: DCT/USO3/05028
CURRENT FILING DATE: 2003-03-11
PRIOR APPLICATION NUMBER: US 60/353,124
PRIOR FILING DATE: 2002-03-11
PRIOR FILING DATE: 2002-06-06
PRIOR FILING DATE: 2002-06-06
PRIOR FILING DATE: 2002-08-09
PRIOR FILING DATE: 2002-08-09
PRIOR FILING DATE: 2002-08-09
PRIOR FILING DATE: 2002-09-05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FEATURE: OTHER INFORMATION: Description of Artificial Sequence: SINA sense region
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LOCATION: (20)...(21)
OTHER INFORMATION: n stands for thymidine
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ORGANISM: Artificial Sequence
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Fosnaugh, Kathy
Jamison, Sharon
Usman, Nassim
Thompson, James
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Best Local Similarity 100.(
Matches 19; Conservative
                                                Thompson, James
                        sman, Nassim
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PCT-US03-05028-450/c
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APPLICANT:
APPLICANT:
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LOCATION: (1). (20)
OTHER INFORMATION: Phosphorothioate 3'-Internucleotide Linkage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  90.5%; Score 19; DB 2; Length 21; 78.9%; Pred. No. 13; Cive 4; Mismatches 0; Indels
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     LoCATION: (20)...(21)
     COTER INFORMATION: n stands for thymidine
PCT-US03-05028-402
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GENERAL INFORMATION:
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: McSwiggen, James
APPLICANT: ChowTira, Bharat
APPLICANT: ChowTira, Bharat
APPLICANT: Pavco, Pamela
APPLICANT: Fosnaugh, Kathy
                                                                                                 PCT-US03-05028-402/c
; Sequence 402, Application PC/TUS0305028
; GENERAL INFORMATION:
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: McSwiggen, James
; APPLICANT: Beigalman, Leonid
; APPLICANT: Chowrira, Bharat
; APPLICANT: Pavco, Pamela
1 CUGAGUUUAAAAGGCACCC 19
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Jamison, Sharon
Usman, Nassim
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Matches 15; Conservative
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APPLICANT:
APPLICANT:
APPLICANT:
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TITLE OF INVENTION: RNA Interference Mediated Inhibition of Gene Expression Using TITLE OF INVENTION: Short Interfering Nucleic Acid (sinA)
TITLE OF INVENTION: Short Interfering Nucleic Acid (sinA)
FILE REPERENCE: 400/104 (MBHB 03-198)
CURRENT APPLICATION NUMBER: 2003-02-20
CURRENT FILING DATE: 2003-02-20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PANENKEY: misc feature
LOCATION: (21) 7. (21)
OTHER INFORMATION: 3'-3 attached terminal deoxyabasic moeity
     LOCATION: (1)..(1)
OTHER INFORMATION: 5'-3 attached terminal deoxyabasic moeity
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Pred. No. 13;
Mismatches 0
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OTHER INFORMATION: n stands for thymidine
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PRIOR PEDILICATION NUMBER: US 60/363,124
PRIOR PEDILICATION NUMBER: US 60/363,124
PRIOR PEDILICATION NUMBER: US 60/368,580
PRIOR FILING DATE: 2002-02-02
PRIOR FILING DATE: 2002-06-06
PRIOR PELING DATE: 2002-06-06
PRIOR PELING DATE: 2002-06-09
PRIOR PELING DATE: 2002-06-29
PRIOR PELING DATE: 2002-09-05
PRIOR PELING DATE: 2002-09-05
PRIOR PELING DATE: 2002-09-09
PRIOR PELING DATE: 2002-09-09
PRIOR PELING DATE: 2002-09-09
PRIOR PELING DATE: 2003-09-09
PRIOR PELING DATE: 2003-09-09
PRIOR PELING DATE: 2003-09-09
PRIOR PELING DATE: 2003-01-15
NUMBER OF SEQ ID NOS: 932
SEQ ID NO 456
                                                                                                                                                                                                        NAME/KEY: misc feature
LOCATION: (15)...(15)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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OTHER INFORMATION: 2'-deoxy-2'-fluoro
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90.5%; Scc
Best Local Similarity 100.0%; Pr
Matches 19; Conservative 0;
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Jamison, Sharon
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                                                                                            NAME/KEY: misc feature
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TITLE OF INVENTION: RNA Interference Mediated Inhibition of Gene Expression Using
TITLE OF INVENTION: Short Interfering Nucleic Acid (sina)
FILE REFERENCE: 400/104 (MBHB 03-198)
CURRENT APPLICATION NUMBER: PCT/US03/05028
CURRENT FILING DATE: 2002 0200
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LOCATION: (20) ... (21)
OTHER INFORMATION: n stands for thymidine
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PRIOR APPLICATION NUMBER: US 60/363,124
PRIOR PELING DATE: 2002-03-11
PRIOR PELING DATE: 2002-03-11
PRIOR PELING DATE: 2002-02-05
PRIOR PELING DATE: 2002-02-06
PRIOR PELING DATE: 2002-06-06
PRIOR PELING DATE: 2002-06-06
PRIOR PELING DATE: 2002-06-06
PRIOR PELING DATE: 2002-06-05
PRIOR PELING DATE: 2002-09-05
PRIOR PELING DATE: 2002-09-05
PRIOR PELING DATE: 2002-09-05
PRIOR PELING DATE: 2002-09-09
PRIOR PELING DATE: 2003-01-15
PRIOR APPLICATION NUMBER: US 60/409,293 PRIOR FLING DATE: 2002-09-09 PRIOR APPLICATION NUMBER: US 60/440,129 PRIOR FILING DATE: 2003-01-15
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LOCATION: (1). (2)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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                                                                                                              NUMBER OF SEQ ID NOS: 932
SOFTWARE: Patentin version 3.2
SEQ ID NO 450
EBNCTH: 21
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SOFWARE: PatentIn version 3.2
SEQ ID NO 453
LENGTH: 21
                                                                                                                                                                                                                                  TYPE: RNA
ORGANISM: Artificial Sequence
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ORGANISM: Artificial Sequence
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Matches 15; Conservative
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Gaps

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ATTLE OF INVENTION: RNA Interference Mediated Inhibition of Gene Expression Using TITLE OF INVENTION: RNA Interfering Nucleic Acid (sinA)
FILE REFERENCE: 400/104 (MBHB 03-198)
GURRENT APPLICATION NUMBER: PCT/US03/05028
CURRENT FILING DATE: 2003-02-20
PRIOR APPLICATION NUMBER: US 60/363,124
PRIOR PLING DATE: 2002-03-11
PRIOR FILING DATE: 2002-03-11
PRIOR FILING DATE: 2002-02-20
PRIOR FILING DATE: US 60/386,782
PRIOR FILING DATE: US 60/386,782
PRIOR PLING DATE: 2002-06-06
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                                                                                               LOCATION: (1). [(1)
OTHER INFORMATION: 5'-3 attached terminal deoxyabasic moeity
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; OTHER INFORMATION: 3'-3 attached terminal deoxyabasic moeity
PCT-USO3-05028-459
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100.0%; Pred. No. 13;
ive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME/KEY: misc_feature
LCGATION: (20)...(21)
OTHER INFORMATION: n stands for thymidine
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       90.5%; Score 19; 100.0%; Pred. No.
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LOCATION: (15)...(15)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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LOCATION: (1)..(2)

STHER INFORMATION: 2'-deoxy-2'-fluoro
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OTHER INFORMATION: 2'-deoxy-2'-fluoro
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OTHER INFORMATION: 2'-deoxy-2'-fluoro
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APPLICANT: McSwiggen, James
APPLICANT: Beigelman, Leonid
APPLICANT: Chowrira, Bharat
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Fosnaugh, Kathy
Jamison, Sharon
Usman, Nassim
Thompson, James
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LOCATION: (9)...(14)
OTHER INFORMATION: 2'-deoxy
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LOCATION: (16)...(16)
OTHER INFORMATION: 2'-deoxy
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OTHER INFORMATION: 2'-deoxy
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                                                                       NAME/KEY: misc_feature
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APPLICANT: PawCo, Pamelia
APPLICANT: Jamison, Sharon
APPLICANT: Jamison, Sharon
APPLICANT: Jamison, Sharon
APPLICANT: Userian, Nassim
APPLICANT: Userian, Names
TITLE OF INVENTION: Short Interfering Nucleic Acid (siNA)
TITLE OF INVENTION: Short Interfering Nucleic Acid (siNA)
FILE OF INVENTION: Short Interfering Number: US 60/363,124
PRIOR APPLICATION NUMBER: US 60/353,124
PRIOR APPLICATION NUMBER: US 60/366,782
PRIOR PILING DATE: 2002-06-06
PRIOR FILING DATE: 2002-09-09
PRIOR FILING DATE: 2002-09-09
PRIOR APPLICATION NUMBER: US 60/408,378
PRIOR FILING DATE: 2002-09-09
PRIOR FILING DATE: 2003-09-09
PRIOR FILING DATE: 2003-09-09
PRIOR FILING DATE: 2003-09-09
PRIOR FILING DATE: 2003-01-15
NUMBER OF SEQ ID NOS: 932
SOFFWARE PATENTING PATE: PatentIn version 3.2
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NAME/KEY: misc_feature
LOCATION: (20)...(20)
CHER INFORMATION: Phosphorothioate 3'-Internucleotide Linkage
PCT-US03-05028-456
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Pred. No. 13;
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LOCATION: (20) ... (21)
OTHER INFORMATION: n stands for thymidine
                                                                                                                 NAME/KEY: misc feature
LOCATION: (6). (11)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                                                                                                                                                      NAME/KEY: misc feature
LOCATION: (15)...(17)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                               LOCATION: (4). (4)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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Best Local Similarity 78.9
Matches 15; Conservative
                     misc_feature
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LENGTH: 21
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OTHER INFORMATION: Description of Artificial Sequence: siNA antisense region
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NAME/KEY: misc_feature
LOCATION: (20)..(21)
OTHER INFORMATION: n stands for thymidine
PRIOR APPLICATION NUMBER: US 60/406,784
PRIOR FILING DATE: 2002-08-29
PRIOR PILING DATE: 2002-08-29
PRIOR PILING DATE: 2002-09-05
PRIOR PILING DATE: 2002-09-09
PRIOR FILING DATE: 2002-09-09
PRIOR FILING DATE: 2003-09-15
PRIOR FILING DATE: 2003-01-15
NUMBER OF SEQ ID NOS: 932
SOFTWARE: PATCHIN VERSION 3.2
SEQ ID NO 462
LENGTH: 21
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NAME/KEY: misc_feature
LOCATION: (4)...(4)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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LOCATION: (15)...(17)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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NAME/KRX: misc feature
LOCATION: (6). (11)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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19 CTGAGTTTAAAAGGCACCC 1
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ORGANISM: Artificial Sequence
FEATURE:
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Search completed: August 18, 2005, 23:33:06 Job time : 3830 secs

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1: /cgn2_6/ptodata/1/pubpna/USO7_PUBCOMB.seq:*
2: /cgn2_6/ptodata/1/pubpna/USO6_NEW PUB.seq:*
3: /cgn2_6/ptodata/1/pubpna/USO6_NEW PUB.seq:*
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6: /cgn2_6/ptodata/1/pubpna/USO6_NEW PUB.seq:*
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7: /cgn2_6/ptodata/1/pubpna/USO9_PUBCOMB.seq:*
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15: /cgn2_6/ptodata/1/pubpna/USO0_PUBCOMB.seq:*
16: /cgn2_6/ptodata/1/pubpna/USO0_PUBCOMB.seq:*
16: /cgn2_6/ptodata/1/pubpna/USO0_PUBCOMB.seq:*
17: /cgn2_6/ptodata/1/pubpna/USO0_PUBCOMB.seq:*
18: /cgn2_6/ptodata/1/pubpna/USO0_PUBCOMB.seq:*
19: /cgn2_6/ptodata/1/pubpna/USO0_PUBCOMB.seq:*
10: /cgn2_6/ptodata/1/pubpna/USOO_PUBCOMB.seq:*
10: /cgn2_6
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/cgn2_6/ptodata/1/pubpna/US11 NEW PUB.seq:*
/cgn2_6/ptodata/1/pubpna/US60_NEW_PUB.seq:*
/cgn2_6/ptodata/1/pubpna/US60_PUBCOMB.seq:*
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                                              GenCore version 5.1.6
Copyright (c) 1993 - 2005 Compugen Ltd.
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Maximum Match 100%
Listing first 45 summaries
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21
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Perfect score:
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2313, 2313, 2313, 2313, 2092,	
Description Sequence Sequence Sequence Sequence Sequence Sequence	Sequence
US-10-665-951-2313 US-10-665-951-2313 US-10-758-155-2313 US-10-814-076-2313 US-10-665-951-2092 US-10-665-951-2096	US-10-665-951-2100
DB 119 22 22 22 19	19
Query Match Length DB 90.5 19 21 90.5 19 22 90.5 19 22 90.5 19 22	21
1	90.5
Score 199-199-199-199-199-199-199-199-199-199	19
ult No. 1 1 3 3 4	7
Result No.	

Sequence 2104, Sequence 2149, Sequence 2155, Sequence 2155, Sequence 2156, Sequence 2168, Sequence 2186, Sequence 2186, Sequence 2218, Sequence 2218, Sequence 2220, Sequence 2221, Sequence 2221, Sequence 2224, Sequence 2224, Sequence 2224, Sequence 2224, Sequence 2224,	Sequence 301, App Sequence 305, App Sequence 315, App Sequence 621, App Sequence 622, App Sequence 622, App Sequence 315, App Sequence 622, App Sequence 622, App Sequence 622, App Sequence 301, App Sequence 301, App Sequence 301, App Sequence 622, App Sequence 622, App Sequence 2092, App Sequence 2092, App Sequence 2092, App Sequence 2100, App Sequence 2109, App Sequence	nhibition of Vascular Endothelial Endothelial Growth Factor Receptor t Interfering Nucleic Acid (siNA)
	US-10-444-853A-301 US-10-444-853A-305 US-10-444-853A-621 US-10-444-853A-622 US-10-757-803-301 US-10-757-803-301 US-10-757-803-622 US-10-757-803-622 US-10-757-803-622 US-10-757-803-622 US-10-757-803-622 US-10-757-803-622 US-10-758-906-622 US-10-826-966-305 US-10-826-966-305 US-10-826-966-622 US-10-826-966-622 US-10-826-966-622 US-10-826-966-622 US-10-758-155-2104 US-10-758-155-2104 US-10-758-155-2104 US-10-758-155-2104	Mediated I d Vascular Using Shox 1,951 68 68 69 49 47 7674
		cation US/1040138163A1 erapeutics, en, James han, Leonid Panel Leonid Growth Fa Growth Fa Growth Fa Gone Exp. (7131 (MRHER: US 12003-09-18 UMBER: US 12002-07-03 UMBER: US 12002-07-03 UMBER: US 12002-07-03 UMBER: US 12002-11-07 UMBER: ECT/104014 US 12002-11-07 UMBER: US 12002-05-29 UMBER: US 12002-05-29
	199 990 55 1199 99	RESULT 1 1. Carlot 1. Carlot 2313, Application US/10665951 1. Publication No. US20040138163A1 1. GENERAL INFORMATION: 1. APPLICANT: Sirna Therapeutics, Inc. 1. APPLICANT: Beagelman, Leonid 1. APPLICANT: Beagelman, Leonid 1. APPLICANT: Pavco, Pamela 1. TITLE OF INVENTION: RNA Interference 1. TITLE OF INVENTION: Growth Factor an 1. TITLE OF INVENTION: Growth Factor an 1. TITLE OF INVENTION: RNA Interference 1. TITLE OF INVENTION: Growth Factor an 1. TITLE OF INVENTION: WUMBER: US/10/665 1. PRIOR APPLICATION NUMBER: US 60/399, 3 1. PRIOR FILING DATE: 2002-07-03 1. PRIOR APPLICATION NUMBER: US 60/393, 7 1. PRIOR FILING DATE: 2002-07-03 1. PRIOR FILING DATE: 2002-05-29
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NAME/KEY: misc_feature
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GENERAL INFORMATION

GRAPLICANT: Sirna Therapeutics, Inc.

APPLICANT: Sirna Therapeutics, Inc.

APPLICANT: Beigelman, Leonid

APPLICANT: Beigelman, Leonid

APPLICANT: Beigelman, Leonid

APPLICANT: Bescentant, Leonid

APPLICANT: Bescentant, Leonid

APPLICANT: Bescentant, Leonid

APPLICANT: Bescentant, Leonid

TITLE OF INVENTION: Grawth Factor and Vascular Endothelial

TITLE OF INVENTION: Graw Expression Using Short Interfering Nucleic Acid (sinA)

FILLE REFERENCE: 400/141 (MHB02742-N)

CURRENT APPLICATION NUMBER: US 10/665,951

PRIOR PLILNG DATE: 2003-09-18

PRIOR PLILNG DATE: 2003-09-18

PRIOR APPLICATION NUMBER: US 60/399,348

PRIOR PLILNG DATE: 2003-07-29

PRIOR PLILNG DATE: 2002-07-29

PRIOR PLILNG DATE: 2002-07-29

PRIOR APPLICATION NUMBER: US 60/399,348

PRIOR PLILNG DATE: 2002-07-29

PRIOR APPLICATION NUMBER: US 60/399,348

PRIOR PLILNG DATE: 2002-11-04

PRIOR PLILNG DATE: 2002-11-04

PRIOR PLILNG DATE: 2002-11-27

PRIOR PLILNG DATE: 2002-05-29

PRIOR PLILNG DATE: 2002-05-29

PRIOR APPLICATION NUMBER: US 60/336,124

PRIOR PLILNG DATE: 2002-05-29

PRIOR APPLICATION NUMBER: US 60/336,124

PRIOR PLILNG DATE: 2002-01-27

PRIOR APPLICATION NUMBER: US 60/336,124

PRIOR PLILNG DATE: 2002-01-27

PRIOR APPLICATION NUMBER: US 60/336,124

PRIOR PLILNG DATE: 2002-01-127

PRIOR APPLICATION NUMBER: US 60/336,124

PRIOR PLILNG DATE: 2002-01-127

PRIOR PLILNG DATE: 2002-01-127

PRIOR PLILNG DATE: 2002-01-127

PRIOR APPLICATION NUMBER: US 60/336,124

PRIOR PLILNG DATE: 2002-01-127

PRIOR PLILNG DATE: 2002-01-127
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       OTHER INFORMATION: Description of Artificial Sequence: sinh antisense region
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                                  PRIOR APPLICATION NUMBER: US 60/363,124
PRIOR FILING DATE: 2002-03-11
PRIOR APPLICATION NUMBER: US 60/386,782
PRIOR FILING DATE: 2002-06-06
REmaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 2455
SOFTWARE: PatentIn version 3.2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Remaining Prior Application data removed - See File Wrapper or PALM. NUMBER OF SEQ ID NOS: 2751
SOFTWARE: PatentIn version 3.3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LOCATION: (20)...(20)
OTHER INFORMATION: Phosphorothioate 3'-Internucleotide Linkage
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME/KEY: misc_feature

) LOCATION: (20)...(20)

) CTHER INFORMATION: X stands for nitroindole universal base

US-10-665-951-2313
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Publication No. US20050075304A1
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  78.9%;
                                                                                                                                                                                                                                                                                                                                                                            TYPE: RNA
ORGANISM: Artificial Sequence
FILING DATE: 2002-02-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity 78.9
Matches 15, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FEATURE:
NAME/KEY: misc_feature
                                                                                                                                                                                                                                                                                                   SEQ ID NO 2313
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LENGTH: 19
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siNA antisense region
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          LOCATION: (20)...(20)
OTHER INFORMATION: Phosphorothioate 3'.-Internucleotide Linkage
                                                                              Length 19;
                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                           OTHER INFORMATION: Description of Artificial Sequence:
                                                                              Score 19; DB 2
Pred. No. 4.2;
4; Mismatches
                                                                                                       1 CUGAGUUUAAAAGGCACCC 19
                                                                                                              CTGAGTTTAAAAGGCACCC 1
                                                                               90.5%;
                                                                                                                                                                                                                                                                                                                                                                                         TYPE: RNA
ORGANISM: Artificial Sequence
ORGANISM: Artificial Sequence
                                                                              Query Match
Best Local Similarity 78.93
Matches 15; Conservative
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CUGAGUUUAAAAGGCACCC 19
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         19 CTGAGTTTAAAAGGCACCC 1
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ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME/KEY: misc_feature
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GRAPLICANT: Sirna Therapeutics, Inc.

APPLICANT: Sirna Therapeutics, Inc.

APPLICANT: Beigelman, Leonid

TITLE OF INVENTION: Growth Factor and Vascular Endothelial

CURRENT APPLICATION NUMBER: US 10/764,957

PRIOR APPLICATION NUMBER: US 10/75,803

PRIOR FILING DATE: 2004-01-12

PRIOR PILING DATE: 2004-01-12

PRIOR APPLICATION NUMBER: US 10/750,448

PRIOR PILING DATE: 2003-11-24

PRIOR FILING DATE: 2003-11-24

PRIOR FILING DATE: 2003-10-23

PRIOR FILING DATE: 2003-10-23

PRIOR FILING DATE: 2003-09-23

PRIOR FILING DATE: 2003-09-23

PRIOR FILING DATE: 2003-09-18

PRIOR PILING DATE: 2003-09-18
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                                                                                                                                                                                        Gaps
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SOFTWARE: PatentIn version 3.3
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LOCATION: (20)<sup>7</sup>.(20)
OTHER INFORMATION: Phosphorothioate 3'-Internucleotide Linkage
i LOCATION: (20)..(20)
i OTHER INFORMATION: X stands for nitroindole universal base
US-10-831-620-2313
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 90.5%; Score 19; DB 22; Length 19; Best Local Similarity 78.9%; Pred. No. 4.2; Matches 15; Conservative 4; Mismatches 0; Indels
                                                                                                                             Length 19
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1 LOCATION: (20)...(20)

2 CTHER INFORMATION: X stands for nitroindole universal base

US-10-844-076-2213
                                                                                                                                                                                     0; Indels
                                                                                                                          Score 19; DB 22;
Pred. No. 4.2;
                                                                                                                                                                                     4; Mismatches .
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Publication No. US20050171039A1
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                19 CTGAGTTTAAAAGGCACCC 1
                                                                                                                          90.5%;
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ORGANISM: Artificial Sequence
                                                                                                                   Query Match
Best Local Similarity 78.9
Matches 15, Conservative
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US-10-844-076-2313/c
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Sequence 2092, Application US/1065951
Sequence 2092, Application US/1065951
Sequence 2092, Application US/1065951
GRNERAL INFORMATION:
MOUSTOORMATION:
MAPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Mexigagen, James
APPLICANT: Moust Demoid
TITLE OF INVENTION: RNA Interference Mediated Inhibition of Vascular Endothelial
TITLE OF INVENTION: Gene Expression Using Short Interfering Nucleic Acid (siNA)
FILE REPRENCE: 400/131 (MBHB02-14-P)
CURRENT PELICATION NUMBER: US/10/65,951
CURRENT PELICATION NUMBER: PCT/US 03/0502
PRIOR FILING DATE: 2003-09-18
PRIOR FILING DATE: 2003-09-18
PRIOR FILING DATE: 2003-09-18
PRIOR FILING DATE: 2002-07-20
PRIOR PELICATION NUMBER: US 60/339,348
PRIOR FILING DATE: 2002-07-03
PRIOR FILING DATE: 2002-07-03
PRIOR FILING DATE: 2002-07-03
PRIOR FILING DATE: 2002-07-03
PRIOR FILING DATE: 2002-01-04
PRIOR PLILING DATE: 2002-01-04
PRIOR FILING DATE: 2002-01-05
PRIOR P
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i OTHER INFORMATION: Phosphorothioate 3'-Internucleotide Linkage
US-10-665-951-2092
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LOCATION: (1). (5) OTHER INFORMATION: Phosphorothioate 3'-Internucleotide Linkage
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OTHER INFORMATION: n stands for thymidine
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; Sequence 2096, Application US/10665951
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TITLE OF INVENTION: RNA Interference Mediated Inhibition of Vascular Endothelial
TITLE OF INVENTION: Growth Factor and Vascular Endothelial Growth Factor Receptor
FILE REFERENCE: 400/131 (MBHB02-742-F)
CURRENT PAPLICATION NUMBER: US/10/665,951
CURRENT APPLICATION NUMBER: US/10/664,668
PRIOR FILING DATE: 2003-09-18
PRIOR FILING DATE: 2003-09-18
PRIOR FILING DATE: 2003-02-0
PRIOR FILING DATE: 2003-02-0
PRIOR FILING DATE: 2003-02-0
PRIOR FILING DATE: 2002-07-29
PRIOR PLICATION NUMBER: US 60/399,348
PRIOR PLICATION NUMBER: US 60/399,796
PRIOR FILING DATE: 2002-07-03
PRIOR FILING DATE: 2002-11-04
PRIOR FILING DATE: 2002-11-04
PRIOR PLICATION NUMBER: US 10/267,949
PRIOR PLICATION NUMBER: US 60/389,580
PRIOR PLICATION NUMBER: PCT/US 02/17674
PRIOR PLICATION NUMBER: US 60/389,580
PRIOR PLICATION NUMBER: US 60/389,580
PRIOR PLICATION NUMBER: US 60/389,782
PRIOR PLICATION NUMBER: US 60/386,782
PRIOR PLICATION NUMBER: US 60/386,782
PRIOR PLILING DATE: 2002-03-01
PRIOR PLILING DATE: 2002-03-06-06
PRIOR PLILING DATE: 2002-03-11
PRIOR PRILICATION NUMBER: US 60/386,782
PRIOR PRILING DATE: 2002-03-06-06
PRIOR PLILING DATE: 200
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LOCATION: (1)._(2)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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NAME/KEY: misc_feature
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LOCATION: (6)..(8)
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LOCATION: (15)..(15)
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                                                                              GENERAL INFORMATION.

APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Bejeslman, Leonid
TITLE OF INVENTION: Growth Factor and Vascular Endothelial
TITLE OF INVENTION: Only 10/65,951
CURRENT FILING DATE: 2003-09-18
FRIOR RELIGATION NUMBER: US 60/399,348
FRIOR FILING DATE: 2002-07-3
FRIOR APPLICATION NUMBER: US 10/306,747
FRIOR APPLICATION NUMBER: US 10/306,747
FRIOR FILING DATE: 2002-11-04
FRIOR FILING DATE: 2002-01-16
FRIOR FILING DATE: 2002-01-16
FRIOR FILING DATE: 2002-01-16
FRIOR APPLICATION NUMBER: US 60/386,780
FRIOR FILING DATE: 2002-02-10
FRIOR FILING DATE: 2002-03-11
FRIOR APPLICATION NUMBER: US 60/386,780
FRIOR FILING DATE: 2002-03-11
FRIOR FILING DATE: 2002-03-11
FRIOR APPLICATION NUMBER: US 60/386,780
FRIOR FILING DATE: 2002-03-11
FRIOR APPLICATION NUMBER: US 60/386,780
FRIOR FILING DATE: 2002-03-11
FRIOR APPLICATION NUMBER: US 60/386,780
FRIOR FILING DATE: 2002-03-11
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i LOCATION: (20)...(20)
i OTHER INFORMATION: Phosphorothioate 3'-Internucleotide Linkage
US-10-665-951-2096
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OTHER INFORMATION: Phosphorothioate 3'-Internucleotide Linkage
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LOCATION: (20)...(21)
OTHER INFORMATION: n stands for thymidine
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ORGANISM: Artificial Sequence
                        Publication No. US20040138163A1
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NAME/KEY: misc_feature
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LOCATION: (1). 7(4)
OTHER INFORMATION: Phosphorothioate 3'-Internucleotide Linkage
                                                                                                                                                                                                                                                                                           LOCATION: (17)...(20)
OTHER INFORMATION: Phosphorothioate 3'-Internucleotide Linkage
                                                                                                                                                                                                                                                                                                                                                                                                       90.5%; Score 19; DB 19; Length 21; ilarity 100.0%; Pred. No. 4.3; Conservative 0; Mismatcht.
                                                                                                                                                                                                                                                                                                                                                            LOCATION: (20)...(21)
OTHER INFORMATION: n stands for thymidine
                                                                                                                                OTHER INFORMATION: 2'-deoxy-2'-fluoro FEATURE:
                                                                                                                                                                                                NAME/KEY: misc feature
LOCATION: (17)...(19)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                                     OTHER INFORMATION: 2'-deoxy-2'-fluoro
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                                                                                                                                                                                                                                                                                                                         FEATURE:
NAME/KEY: misc_feature
                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
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; Sequence 2100, Application US/10665951; Publication No. US20040138163A1; GENERAL INFORMATION:

US-10-665-951-2100

APPLICANT: Sira Therapeutics, Inc.; APPLICANT: Sira Therapeutics, Inc.; APPLICANT: McSwiggen, James; APPLICANT: Beigelman, Leonid; APPLICANT: Pavco, Pamela

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0; Mismatches
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1 LOCATION: (20)...(21)

. OTHER INFORMATION: n stands for thymidine

US-10-665-951-2149
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Publication No. US20040138163A1
GENERAL INFORMATION:
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SEQ ID NO 2149
LENGTH: 21
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Best Local Similarity
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US-10-665-951-2152/c
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                                                                                                                               GENERAL INFORMATION:
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: McSwiggen, James
APPLICANT: Beigelman, Leonid
APPLICANT: Beigelman, Leonid
APPLICANT: Pavoo, Pamela
ITILE OF INVENTION: Growth Factor and Vascular Endothelial
ITILE OF INVENTION: Growth Factor and Vascular Endothelial Growth Factor Receptor
ITILE OF INVENTION: Growth Factor and Vascular Endothelial Growth Factor and Vascular Endothelial
ITILE OF INVENTION: Growth Factor and Vascular Endothelial Growth Factor Receptor
ITILE OF INVENTION: Growth Factor and Vascular Endothelial
ITILE OF INVENTION: Growth Factor and Vascular Endothelial
ITILE OF INVENTION: Growth Factor and Vascular Endothelial
ITILE OF INVENTION WHERE: US 10/665,951
CURRENT APPLICATION WHERE: US 60/399,348
FRIOR APPLICATION WHERE: US 60/399,348
FRIOR FILING DATE: 2002-07-29
FRIOR FILING DATE: 2002-07-29
FRIOR FILING DATE: 2002-11-04
FRIOR FILING DATE: 2002-03-11
FRIOR FILING DATE: 2002-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OTHER INFORMATION: Description of Artificial Sequence: siNA antisense region
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LOCATION: (1). (20)
OTHER INFORMATION: Phosphorothioate 3'-Internucleotide Linkage
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; OTHER INFORMATION: n stands for thymidine US-10-665-951-2104
; Sequence 2104, Application US/10665951; Publication No. US20040138163A1; GENERAL INFORMATION:
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Sequence 2149, Application US/10665951
Publication No. US20040138163A1
GENERAL INFORMATION:
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ORGANISM: Artificial Sequence
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LOCATION: (20)..(21)
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TITLE OF INVENTION: RNA Interference Mediated Inhibition of Vascular Endothelial
TITLE OF INVENTION: Growth Factor and Vascular Endothelial Growth Factor Receptor
TITLE OF INVENTION: Gene Expression Using Short Interfering Nucleic Acid (sinA)
FILE REPERENCE: 400/131 (MBHB02-742-F)
CURRENT APPLICATION NUMBER: US/10/665,951
CURRENT FILING DATE: 2003-09-18
PRIOR APPLICATION NUMBER: BT/US 03/65022
PRIOR APPLICATION NUMBER: PT/US 03/65022
PRIOR APPLICATION NUMBER: US 60/399,348
PRIOR FILING DATE: 2002-07-29
PRIOR APPLICATION NUMBER: US 60/393,796
PRIOR FILING DATE: 2002-01-04
PRIOR FILING DATE: 2002-01-04
PRIOR FILING DATE: 2002-01-04
PRIOR FILING DATE: 2002-01-1-27
PRIOR FILING DATE: 2002-01-27
PRIOR FILING DATE: 2002-01-27
PRIOR FILING DATE: 2002-01-27
PRIOR PRIOR FILING DATE: 2002-02-20
PRIOR FILING DATE: 2002-03-11
PRIOR FILING DATE: 2002-06-06
PRIOR PRIOR PRIOR DATE: 2002-06-06
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APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: McSwiggen, James
APPLICANT: McSwiggen, James
APPLICANT: McSwiggen, James
APPLICANT: Belgelman, Leonid
APPLICANT: Bavco, Pamela
TITLE OF INVENTION: Row Interference Mediated Inhibition of Vascular Endothelial
TITLE OF INVENTION: Growth Factor and Vascular Endothelial Growth Factor Receptor
TITLE OF INVENTION: Growth Factor and Vascular Endothelial
TITLE OF INVENTION: Gene Expression Using Short Interfering Nucleic Acid (sinA)
FILE REFERENCE: 400/131 (MBHB02-742-F)
CURRENT FILING DATE: 2003-09-18
PRIOR FILING DATE: 2003-09-18
PRIOR APPLICATION NUMBER: US 10/664,668
PRIOR PELING DATE: 2003-09-18
PRIOR PILING DATE: 2003-09-00-18
PRIOR FILING DATE: 2003-02-00
PRIOR APPLICATION NUMBER: US 60/399,348
PRIOR FILING DATE: 2000-07-03
PRIOR FILING DATE: 2000-07-03
PRIOR FILING DATE: 2000-07-03
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##FILLE OF INVENTION: Growth Factor and Vascular Endothelial
TITLE OF INVENTION: Growth Factor and Vascular Endothelial Growth Factor Receptor
TITLE OF INVENTION: Growth Factor and Vascular Endothelial Growth Factor Receptor
TITLE OF INVENTION: Gene Expression Using Short Interfering Nucleic Acid (siNA)
TITLE OF INVENTION: Gene Expression Using Short Interfering Nucleic Acid (siNA)
FILE REFERENCE: 400/131 (MBHB02-742-F)
CURRENT APPLICATION NUMBER: US/10/665,951
CURRENT PAPLICATION NUMBER: US 10/664,668
PRIOR FILING DATE: 2003-09-18
PRIOR FILING DATE: 2003-02-20
PRIOR APPLICATION NUMBER: US 60/399,348
PRIOR FILING DATE: 2002-07-03
PRIOR PILING DATE: 2002-07-03
PRIOR FILING DATE: 2002-07-03
PRIOR FILING DATE: 2002-11-04
PRIOR FILING DATE: 2002-11-27
PRIOR FILING DATE: 2002-11-27
                                                                                                                                                                                                                                                                 OTHER INFORMATION: Description of Artificial Sequence: siNA sense region
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LOCATION: (1)...(1)
OTHER INFORMATION: 5'-3 attached terminal deoxyabasic moeity
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OTHER INFORMATION: n stands for thymidine
                                                                                                                                                                                                                                                                                                                                             NAME/KEY: misc feature
LOCATION: (1)..(2)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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OTHER INFORMATION: 2'-deoxy-2'-fluoro
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OTHER INFORMATION: 2'-deoxy-2'-fluoro
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Publication No. US20040138163A1
GENERAL INFORMATION:
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                                                                                                                                           TYPE: RNA
ORGANISM: Artificial Sequence
FEATURE:
                     SOFTWARE: PatentIn version 3.2
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NAME/KEY: misc_feature
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NAME/KEY: misc_feature
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Best Local Similarity
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Sequence 2155, Application US/10665951

Sequence 2155, Application US/10665951

GENERAL INFORMATION:

APPLICANT: SITUAT Therapeutics, Inc.

APPLICANT: Beigelman, Leonid

TITLE OF INVENTION: Gene Expression Using Short Interfering Nucleic Acid (sinA)

TITLE OF INVENTION: Gene Expression Using Short Interfering Nucleic Acid (sinA)

TITLE OF INVENTION: Gene Expression Using Short Interfering Nucleic Acid (sinA)

TITLE OF INVENTION: Gene Expression US/10/665,951

CURRENT APPLICANTON NUMBER: US/10/665,951

CURRENT PLING DATE: 2003-09-18

PRIOR PLING DATE: 2003-09-22

PRIOR APPLICATION NUMBER: US 60/393,796

PRIOR APPLICATION NUMBER: US 60/393,249

PRIOR APPLICATION NUMBER: US 60/393,249

PRIOR APPLICATION NUMBER: US 60/389,580

PRIOR APPLICATION NUMBER: US 60/389,580

PRIOR APPLICATION NUMBER: US 60/386,780

PRIOR APPLICATION NUMBER: US 60/386,780

PRIOR APPLICATION NUMBER: US 60/386,780

PRIOR APPLICATION NUMBER: US 60/386,780
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PRIOR APPLICATION NUMBER: US 10/287,949
PRIOR FILING DATE: 2002-11-04
PRIOR PILING DATE: 2002-11-04
PRIOR PILING DATE: 2002-11-27
PRIOR PILING DATE: 2002-05-29
PRIOR FILING DATE: 2002-05-29
PRIOR PILING DATE: 2002-05-29
PRIOR PILING DATE: 2002-02-20
PRIOR PILING DATE: 2002-02-20
PRIOR PILING DATE: 2002-03-11
PRIOR PILING DATE: 2002-03-11
PRIOR PILING DATE: 2002-03-11
PRIOR PILING DATE: 2002-06-06
PRIOR PILING DATE: 2002-06
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Pred. No. 4.3;
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4; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; NAME/KEY: misc_feature
; LOCATION: (20)...(21)
... OTHER INFORMATION: n stands for thymidine
US-10-665-951-2152
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APPLICATION NUMBER: US 60/386,782
FILING DATE: 2002-06-06
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ORGANISM: Artificial Sequence
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78.9%;
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Best Local Similarity 78.9°
Matches 15, Conservative
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OTHER INFORMATION: Description of Artificial Sequence: sinA sense region
PRIOR FILING DATE: 2002-11-27
PRIOR APPLICATION NUMBER: PCT/US 02/17674
PRIOR PLICATION NUMBER: PCT/US 02/17674
PRIOR FILING DATE: 2002-05-29
PRIOR FILING DATE: 2002-05-29
PRIOR PLING DATE: 2002-02-20
PRIOR PLING DATE: 2002-03-11
PRIOR PLING DATE: 2002-03-11
PRIOR PLING DATE: 2002-03-11
PRIOR PLING DATE: 2002-03-11
PRIOR PLING DATE: 2002-06-06
Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 2455
SOFTWARE: PatentIn version 3.2
LENGTO.
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LOCATION: (1). [1)
OTHER INFORMATION: 5'-3 attached terminal deoxyabasic moeity
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; LOCATION: (21):.(21)
; OTHER INFORMATION: 3'-3 attached terminal deoxyabasic moeity
US-10-665-951-2161
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4.3;
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OTHER INFORMATION: n stands for thymidine
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LOCATION: (1)..(2)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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LOCATION: (6). (8). (8)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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INFORMATION: 2'-deoxy-2'-fluoro
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OTHER INFORMATION: 2'-deoxy-2'-fluoro
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US-10-665-951-2164/c
; Sequence 2164, Application US/10665951
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ORGANISM: Artificial Sequence
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LOCATION: (9)...(14)
OTHER INFORMATION: 2'-deoxy
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OTHER INFORMATION: 2'-deoxy
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OTHER INFORMATION: 2'-deoxy
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LOCATION: (3)..(5)
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APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: Beigglan, James
APPLICANT: Beigglan, Leonid
APPLICANT: Pavco, Pamela
TITLE OF INVENTION: Growth Factor and Vascular Endothelial
TITLE OF INVENTION: Growth Factor Alone
FILE REFERENCE: 400/131 (MBHBR: US)10/665,951
CURRENT FILING DATE: 2003-09-18
PRIOR APPLICATION NUMBER: US 10/64,668
PRIOR FILING DATE: 2003-09-18
PRIOR FILING DATE: 2003-09-18
PRIOR FILING DATE: 2002-07-29
PRIOR APPLICATION NUMBER: US 60/393,796
PRIOR APPLICATION NUMBER: US 60/393,796
PRIOR APPLICATION NUMBER: US 60/393,796
PRIOR FILING DATE: 2002-07-03
PRIOR APPLICATION NUMBER: US 60/393,796
PRIOR FILING DATE: 2002-07-03
PRIOR FILING DATE: 2002-07-03
PRIOR FILING DATE: 2002-11-04
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                                                            PRIOR FILLING DATE: 2002-05-29
PRIOR APPLICATION NUMBER: US 60/358,580
PRIOR APPLICATION NUMBER: US 60/358,580
PRIOR FILING DATE: 2002-03-10
PRIOR PLING DATE: 2002-03-11
PRIOR FILING DATE: 2002-06-06
Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 2455
SCOTWARE: Patentin version 3.2
SEQ ID NO 2158
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// LOCATTON: (20)...(20)
// OTHER INFORMATION: Phosphorothioate 3'-Internucleotide Linkage
US-10-665-951-2158
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Pred. No. 4.3;
4; Mismatches 0; Indels
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LOCATION: (20)..(21)
OTHER INFORMATION: n stands for thymidine
                                                 APPLICATION NUMBER: PCT/US 02/17674
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NAME/KEY: misc_feature
LOCATION: (4)...(4)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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OTHER INFORMATION: 2'-deoxy-2'-fluoro
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Publication No. US20040138163A1
GENERAL INFORMATION:
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ORGANISM: Artificial Sequence
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Best Local Similarity 78.9
Matches 15; Conservative
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Query Match
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                                                                                                                                                             ## APPLICANT: Bavco, Pamela
| TITLE OF INVENTION: RNA Interference Mediated Inhibition of Vascular Endothelial
| TITLE OF INVENTION: RNA Interference Mediated Inhibition of Vascular Endothelial
| TITLE OF INVENTION: Growth Factor and Vascular Indothelial Growth Factor Receptor ITILE OF INVENTION: Gene Expression Using Short Interfering Nucleic Acid (BinA)
| TITLE OF INVENTION: Gene Expression Using Short Interfering Nucleic Acid (BinA)
| CURRENT APPLICATION NUMBER: US/10/665,951
| CURRENT PLINGS DATE: 2003-09-18
| PRIOR APPLICATION NUMBER: US 60/339,348
| PRIOR PRILING DATE: 2003-00-18
| PRIOR PELLON NUMBER: US 60/339,348
| PRIOR PELLON NUMBER: US 60/339,348
| PRIOR PELLON NUMBER: US 60/339,348
| PRIOR PELLON NUMBER: US 10/287,949
| PRIOR PELLON NUMBER: US 10/287,949
| PRIOR PELLON NUMBER: US 10/287,949
| PRIOR APPLICATION NUMBER: US 60/338,580
| PRIOR PELLON NUMBER: US 60/338,580
| PRIOR PELLON NUMBER: US 60/358,580
| PRIOR PELLON NUMBER: US 60/358,580
| PRIOR PELLON NUMBER: US 60/358,580
| PRIOR PELLON NUMBER: US 60/388,580
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2 OTHER INFORMATION: Phosphorothioate 3'-Internucleotide Linkage
US-10-665-951-2164
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OTHER INFORMATION: n stands for thymidine
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LOCATION: (6)._(11)
OTHER INFORMATION: 2'-deoxy-2'-fluoro
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OTHER INFORMATION: 2'-deoxy-2'-fluoro
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OTHER INFORMATION: 2'-deoxy-2'-fluoro
                                                                           APPLICANT: Sirna Therapeutics, Inc.
APPLICANT: McSwiggen, James
APPLICANT: Beigelman, Leonid
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ORGANISM: Artificial Sequence
Publication No. US20040138163A1
GENERAL INFORMATION:
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LOCATION: (4)...(4)
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Best Local Similarity
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US-10-665-591-2185
Publication No. USC0040138163A1
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NAME/KRY: misc feature
LOCATION: (1)..(1)
OTHER INFORMATION: 5'-3 attached terminal deoxyabasic moeity
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OTHER INFORMATION: 3'-3 attached terminal deoxyabasic moeity
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OTHER INFORMATION: n stands for thymidine FEATURE:
NAME/KEY: misc_feature
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NAME/KEY: misc_feature
LOCATION: (20)..(21)
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Matches 19; Conserv
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APPLICANT: Bavco, Pamela
APPLICANT: Bavco, Pamela
APPLICANT: Barinchcomb, Dan T
APPLICANT: Stinchcomb, Dan T
APPLICANT: Stinchcomb, Dan T
APPLICANT: Bacobedo, Jaime
TITLE OF INVENTION: TREATMENT OF DISEASES OR
TITLE OF INVENTION: CONDITIONS RELATED TO LEVELS
TITLE OF INVENTION: OF VASCULAR ENDOTHELIAL
TITLE OF SEQUENCES: 8502
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CITY: Los Angeles
STATE: California
COUNTRY: U.S.A.
ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US/08/584,040
FILING DATE: January 11, 1996
CLASSIPICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/005,974
FILING DATE: October 26, 1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 1494, Application US/08584040 Patent No. 6346398 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFRENCE/DOCKET NUMBER: 218/064
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEPAX: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
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SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
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7242, Ap
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30689, A
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Sequence 39, Appl
Sequence 24, Appl
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                                                                                                       August 18, 2005, 21:39:47 ; Search time 124 Seconds (without alignments) 277.111 Million cell updates/sec
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/cgn2_6/ptodata/1/ina/6A_COMB.seq:*
/cgn2_6/ptodata/1/ina/6B_COMB.seq:*
/cgn2_6/ptodata/1/ina/PCTUS_COMB.seq:*
/cgn2_6/ptodata/1/ina/PCTUS_COMB.seq:*
GenCore version 5.1.6
Copyright (c) 1993 - 2005 Compugen Ltd.
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US-09-583-318-24
US-08-584-040-12423
US-08-584-040-12423
US-09-371-7728-38
US-09-685-6648-38
US-09-953-318-25
US-09-584-040-7241
US-09-3741-7728-37
US-09-396-1966-30680
US-09-396-1966-30680
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US-09-396-196G-106340
US-08-687-080-162
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US-09-422-978-9871
US-09-396-196G-44902
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Maximum Match 100%
Listing first 45 summaries
                                                                        nucleic search, using sw model
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Match Length DB
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	34 12 35 12 36 11.8	212	57.1	17 28 17	4 4 4 4	US-09-371-772B-4222 US-08-877-966B-8 US-09-080-044-15 US-09-531-857A-15	Sequence Sequence Sequence	4222, Ap 8, Appli 15, Appl
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ALIGNMENTS

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Best Local Similarity 100.
Matches 17; Conservative
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Best Local Similarity 100.
Matches 17; Conservative
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                                 Score 17; DB 3; Length 17;
Pred. No. 7.6;
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                                                                                                                                                                                                                                                                    APPLICANT: Pavco, Pamela
APPLICANT: Pavco, Pamela
APPLICANT: Pavcomb, Dan T.
APPLICANT: Escinchcomb, Dan T.
APPLICANT: Escobedo, Jaime
TITLE OF INVENTION: METHOD AND REAGENT FOR THE
TITLE OF INVENTION: TREATMENT OF DISEASES OR
TITLE OF INVENTION: GROWTH FACTOR
NUMBER OF SIQUENCES: 8502
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
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                                                                   Mismatches
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STREET: 633 West Fifth Street
STREET: Suite 4700
CITY: Los Angeles
STATE: California
COUNTRY: U.S.A.
ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: Storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM COMPACTION
COMPUTER: IBM COMPACTION
SOFTWARE: WORD PERFECT 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/584,040
FILING DATE: January 11, 1996
CLASSIFICATION DATA:
PRIOR APPLICATION NUMBER: 60/005,974
FILING DATE: OCLOBER 26, 1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REERENCE/DOCKET NUMBER: 218/064
TELEPHONE: (213) 489-1600
TELEPHONE: (213) 955-0440
TELERA: (213) 955-0440
TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 7243:
SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                   Sequence 7243, Application US/08584040; Patent No. 6346398; GENERAL INFORMATION:
                   81.0%; SC.
100.0%; Pre
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100.0%; Pre
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Best Local Similarity 100.
Matches 17; Conservative
                                                                 Conservative
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               Query Match
Best Local Similarity
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US-08-584-040-1494
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RESULT 3

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APPLICANT: Pavco, Pam
APPLICANT: Pavco, Pam
APPLICANT: Pavco, Pam
APPLICANT: Stinchcoub, Dan
APPLICANT: Stinchcoub, Dan
APPLICANT: Stinchcoub, Dan
APPLICANT: Escobedo, Jaime
APPLICANT: Escobedo, Jaime
APPLICANT: Escobedo, Jaime
APPLICANT: Levels of Vascular Endothelial Growth Factor Receptor
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
FILE REFERENCE: MHHBO, 876-3 (237/198)
CURRENT APPLICATION NUMBER: US/09/371,772B
CURRENT APPLICATION NUMBER: US 60/005,974
PRIOR FILING DATE: 1995-10-26
PRIOR FILING DATE: 1995-01-08
NUMBER OF SEQ ID NOS: 14225
SOFTWARE: Patentin version 3.0
SEQ ID NO 39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Pavco, Pam

APPLICANT: Pavco, Pam

APPLICANT: McSwiggen, Jim

APPLICANT: Bacobodo, Jame

APPLICANT: Stinchcount, Dan

APPLICANT: Stinchcount, Dan

APPLICANT: Stinchcount, Dan

TITLE OF INVENTION: Method and Reagent for Treatment of Diseases or Conditions Relate

TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor

TITLE OF INVENTION: WHENER: US/09/685,664B

CURRENT APPLICATION NUMBER: US 60/005,974

PRIOR APPLICATION NUMBER: US 60/005,974

PRIOR APPLICATION NUMBER: US 08/584,040

PRIOR APPLICATION NUMBER: US 09/371,772

PRIOR FILING DATE: 1995-01-026

PRIOR PRILING DATE: 1995-00-08-10

MUMBER: OF SEQ ID NOS: 8231

SEQ ID NO 39
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Patent No. 6818447
GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
Sequence 39, Application US/09371772B
Patent No. 6566127
GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Pavco, Pamela
APPLICANT: Pavco, Pamela
APPLICANT: Pavco, James
APPLICANT: Stinchcomb, Dan T.
APPLICANT: Stinchcomb, Dan T.
APPLICANT: Escobedo, Jaime
TITLE OF INVENTION: METHOD AND REAGENT FOR THE
TITLE OF INVENTION: CONDITIONS RELATED TO LEVELS
TITLE OF INVENTION: OF VASCULAR ENDOTHELIAL
TITLE OF INVENTION: GROWTH FACTOR
NUMBER OF SEQUENCES: 8502
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                      DB 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        76.2%; Score 16; DB 3;
100.0%; Pred. No. 27;
tive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
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COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/584,040
FILING DATE: January 11, 1996
CLASSIFICATION DATA:
APPLICATION NUMBER: 60/005,974
FILING DATE: OCCODER 26, 1995
ATTORNEY/AGENT INPORMATION:
                                                                                                                                                                                                                                                                                                                   76.2%; Score 16;
100.0%; Pred. No.
tive 0; Mismatc
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STATE: California
COUNTRY: U.S.A.
ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; Sequence 7242, Application US/08584040
; Patent No. 6346398
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: Warburg, Richard J. REGISTRATION NUMBER: 32,327 REFERENCE/DOCKET NUMBER: 21 TELECOMMUNICATION: INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 CUGAGUUUAAAAGGCA 16
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TELEFAX: (213) 955-0440
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
                                                                                                                                                                                                                                                                                                                      Query Match 76.2
Best Local Similarity 100.
Matches 16; Conservative
TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                   LENGTH: 17 base pairs
TYPE: mucleic acid
TYPE: TYPE:
TYPE: TYPE: TYPE:
TYPE: TYPE: TYPE:
TYPE: TYPE: TYPE: TYPE:
TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: TYPE: T
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STRANDEDNESS: single
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Best Local Similarity
Matches 16; Conserva
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US-08-584-040-7242
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                                 Sequence 24, Application US/09953318

Patent No. 6710174

GENERAL INFORMATION:
APPLICANT: C. Frank Bennett
APPLICANT: Andrew T. Watt
TITLE OF INVENTION: ANTISENSE MODULATION OF VASCULAR ENDOTHELIAL GROWTH FACTOR RECEPT
TITLE OF INVENTION: EXPRESSION
FILE REFERENCE: RTS-023
CURRENT APPLICATION NUMBER: US/09/953,318
CURRENT FILING DATE: 2001-09-13
NUMBER OF SEQ ID NOS: 154
LENGTH: 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 1493, Application US/08584040
; Sequence 1493, Application US/08584040
; Patent No. 6346399
; Patent No. 6346399
; Patent No. 6346399
; Patent No. 6346399
; Patent Involved No. 6346399
; Patent Strichcomb, Dan T. APPLICANT: Strichcomb, Dan T. TILLE OF INVENTION: TREATMENT OF DISEASES OR TITLE OF INVENTION: OF VASCULAR ENDOTHELIAL TITLE OF INVENTION: GROWTH FACTOR NUMBER OF SEQUENCES: 8502
; CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; OTHER INFORMATION: Antisense Oligonucleotide
US-09-953-318-24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ZIP: 90071-2066
CCMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb MEDIUM TYPE: 3.5" Diskette, 1.44 Mb MEDIUM TYPE: 1.5" Diskette, 1.44 Mb MEDIUM TYPE: 1.5" Diskette, 1.44 Mb MEDIUM TYPE: 1.5" Diskette, 1.44 Mb MEDIUM TYPE: 1.0" Diskette, 1.44 Mb MEDIUM TYPE: 1.0" Diskette, 1.44 Mb MEDIUM SYTHEN P.C. DOS 5.0 SOFTWARE: Word Perfect 5.1 CURRENT APPLICATION DATA:
APPLICATION NUMBER: 60/005,974
FILING DATE: October 26, 1995
ATTORNEY AGENT THORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 218/064
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 CUGAGUUUAAAAGGCAC 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (213) 955-0440
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STATE: California
COUNTRY: U.S.A.
ZIP: 90071-2066
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US-00-953-318-25/c
US-00-953-318-25/c
US-00-953-318-25/c
Batent No. 6710174
Batent No. 6710174
GENERAL INFORMATION:
APPLICANT: Andrew T. Watt
TITLE OF INVENTION: ANTIENSE MODULATION OF VASCULAR ENDOTHELIAL GROWTH FACTOR RECEPTOR TITLE OF INVENTION: EXPRESSION
TITLE OF INVENTION: ANTIENCE: RTS-0232
CURRENT APPLICATION NUMBER: US/09/953,318
CURRENT APPLICATION NUMBER: US/09-13
NUMBER OF SEQ ID NOS: 154
LENGTH: 20
  Gaps
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  Indels
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Patent No. 6346398

GENERAL INFORMATION:
APPLICANT: Pavco, Pamela
APPLICANT: Pavco, Pamela
APPLICANT: Stinchcomb, Dan T.
APPLICANT: Escobedo, Jaime
TITLE OF INVENTION: METHOD AND REAGENT FOR THE
TITLE OF INVENTION: TREATMENT OF DISEASES OR
TITLE OF INVENTION: CONDITIONS RELATED TO LEVELS
TITLE OF INVENTION: GROWTH FACTOR
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; OTHER INFORMATION: Antisense Oligonucleotide US-09-953-318-25
  Mismatches
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OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/584,040
FILING DATE: January 11, 1996
CLASSIFICATION: 514
PRIOR APPLICATION 514
APPLICATION NUMBER: 60/005,974
FILING DATE: October 26, 1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: METHOD AND RETITLE OF INVENTION: TREATMENT OF TITLE OF INVENTION: CONDITIONS RETITLE OF INVENTION: GROWTH FACTOR NUMBER OF SEQUENCES: 8502
CORRESPONDENCE ADDRESS:
STREET: 633 West Fifth Street STREET: SLILE 4700
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: DNA ORGANISM: Artificial Sequence
                                                         1 CUGAGUUUAAAAGGCA 16
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Best Local Similarity 81.2
Matches 13; Conservative
  16; Conservative
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STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY: U.S.A. ZIP: 90071-2066
Matches
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Sequence 38, Application US/0968564B

Patent No. 681847

GENERAL INFORMATION:
APPLICANT: Ribosyme Pharmaceuticals, Inc.
APPLICANT: Pavco. pam
APPLICANT: Roswiggen, Jim
APPLICANT: Escobedo, Jaime
APPLICANT: Escobedo, Jaime
APPLICANT: Escobedo, Jaime
TITLE OF INVENTION: Method and Reagent for Treatment of Diseases or Conditions Relate
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Recptor
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Recptor
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Recptor
CURRENT FILING DATE: 1995-10-26
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR APPLICATION NUMBER: US 09/371,772
PRIOR APPLICATION NUMBER: US 09/371,772
PRIOR APPLICATION NUMBER: US 09/371,772
PRIOR PELING DATE: 1999-08-10
SPRIOR FILING DATE: 1999-08-10
                                                                                                                                                                                                                                                                                                                            APPLICANT: PLOYING THE INTERCOLLEGES, INC.
APPLICANT: MCSWiggen, Jim
APPLICANT: MCSWiggen, Jim
APPLICANT: McSwiggen, Jim
APPLICANT: StinchCondb, Dan
APPLICANT: Escobedo, Jaime
TITLE OF INVENTION: Levels of Vacular Endothelial Growth Factor Receptor
TITLE OF INVENTION: Levels of Vacular Endothelial Growth Factor Receptor
CURRENT APPLICATION NUMBER: US/09/371, 772B
CURRENT APPLICATION NUMBER: US 60/005, 974
PRIOR APPLICATION NUMBER: US 08/584, 040
PRIOR APPLICATION NUMBER: US 08/584, 040
NUMBER OF SEQ ID MOS: 14225
SOFTWARE: Patentin version 3.0
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Pred. No. 27;
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                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc
                                                                                                                                                                                                                     . Sequence 38, Application US/09371772B ; Patent No. 6566127
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     76.2%; Scor.
100.0%; Pre
                               1 CUGAGUUUAAAAGGCA 16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 76.2
Best Local Similarity 100.
Matches 16; Conservative
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ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ORGANISM: Homo sapiens
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Best Local Similarity
                                                                                                                                                                                           US-09-371-772B-38
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APPLICANT: ALCOUNT: ALCOUNTES, MINE.
APPLICANT: McSwiggen, Jim
APPLICANT: McSwiggen, Jim
APPLICANT: Stinchcomb, Dan
APPLICANT: Stinchcomb, Dan
APPLICANT: Stinchcomb, Dan
APPLICANT: Escobedo, Jaime
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions Re
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
FILE REFERENCE: MBHB00, 376-37(337/198)
CURRENT FILING DATE: 1999-08-10
PRIOR PILING DATE: 1995-10-26
PRIOR PILING DATE: 1995-10-26
PRIOR FILING DATE: 1995-10-26
PRIOR FILING DATE: 1995-10-8
NUMBER OF SEQ ID NOS: 14225
SOFTWARE: Patentin version 3.0
SEQ ID NO 37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: ROSWIGGEN, JIM.
APPLICANT: Pavco, Pam
APPLICANT: Strinchond, Dan
APPLICANT: Estinchond, Dan
APPLICANT: Escobedo, Jaime
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Recptor
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Recptor
TITLE REPERENCE: MBHB00-87-K (400/021)
CURRENT APPLICATION NUMBER: US (09/05,974
PRIOR APPLICATION NUMBER: US (005,974
PRIOR APPLICATION NUMBER: US (08/584,040
PRIOR APPLICATION NUMBER: US (09/371,772
PRIOR APPLICATION NUMBER: US (09/371,772
PRIOR FILING DATE: 1999-08-10
                                                                                              Length 17;
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                                                                                              DB 3;
93;
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100.0%; Pred. No. 93;
tive 0; Mismatches
                                                                                              71.4%; Score 15; DB 100.0%; Pred. No. 93; Live 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                    Sequence 37, Application US/09371772B
Patent No. 6566127
GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 37, Application US/09685664B Patent No. 6818447
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQ ID NOS: 8231
SOFTWARE: Patentin version 3.0
SEQ ID NO 37
                                                                                                                                                                                                                                 3 CUGAGUUUAAAAGGC 17
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                                                                                              Query Match
Best Local Similarity 100.0
Matches 15; Conservative
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Matches 15; Conservative
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ORGANISM: Homo sapiens
; STRANDEDNESS: sir
; TOPOLOGY: linear
US-08-584-040-7241
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US-09-371-772B-37
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US-09-685-664B-37
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Pavco, Pamela
APPLICANT: Pavco, Pamela
APPLICANT: Barcheomb, Dan T
APPLICANT: Stinchcomb, Dan T
APPLICANT: Stinchcomb, Dan T
APPLICANT: Escobedo, Jaime
TITLE OF INVENTION: TREATHENT OF DISEASES OR
TITLE OF INVENTION: CONDITIONS RELATED TO LEVELS
TITLE OF INVENTION: OF VASCULAR ENDOTHELIAL
TITLE OF INVENTION: OF VASCULAR ENDOTHELIAL
TITLE OF SEQUENCES: 8502
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STATE: CALLLL.
COUNTRY: U.S.A.
ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: Grorage
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/584,040
PILING DATE: January 11, 1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION N.L.

FILING DATE: January L.

FILING DATE: January L.

PRIOR APPLICATION: 514

PRIOR APPLICATION NUMBER: 60,005,974

FILING DATE: October 26, 1995

ATTORNEY/AGENT INPORMATION:

NAME: Warburg, Richard J.

REFERENCE/DOCKET NUMBER: 32,327

REFERENCE/DOCKET NUMBER: 32,327

REFERENCE/DOCKET NUMBER: 32,327

REFERENCE/DOCKET NUMBER: 32,327

RELEPHONE: (213) 489-1600

TELEPHONE: (213) 955-0440

TELEPHONE: (213) 955-0440

TELEPHONE: CHARACTERISTICS:

LENGTH: 17 base pairs

TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 7241, Application US/08584040 Patent No. 6346398 GENERAL INFORMATION:
               NAME: WATCHING.

REGISTRATION NUMBER: 32,327

REFERENCE/DOCKET NUMBER: 218/
TELEPHONE: (213) 489-1600

TELEPHONE: (213) 955-0440

TELEPK: 67-3510

INFORMATION FOR SEQ ID NO: 1492:
SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
CITY: Los Angeles
STATE: California
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ATTORNEY/AGENT INFORMATION:
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Best Local Similarity 100.
Matches 15; Conservative
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US-08-584-040-1492
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                                                                                                    Query Match 71.4%; Score 15; DB 4; Length 17; Best Local Similarity 100.0%; Pred. No. 93; Matches 15; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                 RESULT 15
US-09-396-196G-30680
; Sequence 30680, Application US/09396196G
; Patent No. 6821724
; GENERAL INFORMATION:
; APPLICANT: Michael Mittmann
APPLICANT: David Mack
; APPLICANT: David Mack
; APPLICANT: David Lockhart
; APPLICANT: David Mack
; TITLE OF INVENTION: Methods of Genetic Analysis
; FILE REFERENCE: 3101.1
; CURRENT FILING DATE: 1999-09-15
; PRIOR FILING DATE: 1998-09-17
; NUMBER OF SEQ ID NOS: 127806
; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 30680
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                                                                                                                                                                                                             3 CUGAGUUUAAAAGGC 17
                                                                                                                                                                                   1 CUGAGUUUAAAAGGC 15
; LENGTH: 17
; TYPE: RNA
; ORGANISM: Homo sapiens
US-09-685-664B-37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: DNA
CORGANISM: Mus musculus
US-09-396-196G-30680
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Search completed: August 18, 2005, 22:27:29 Job time : 125 secs

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1M0537N06

Scoring table:

Searched:

Minimum I Maximum I

Database

Perfect score:

Sequence:

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Run on:

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AZ309241
1M0013106F Mouse 10kb plasmid UUGCIM library Mus musculus genomic clone UUGCIM0013106 F, genomic survey sequence.
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Dunn, D., Aoyaqi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Niederhausern, A. and Wright, D., Weise, R., Tingey, A., von Niederhausern, A. and Wright, D., Weise, R.
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                AZ387817 1M0147B24
AZ584523 1M0389C10
AUZ54095 AUZ54095
AZ789794 2M0037J21
AZ783417 2M0025C07
D19150 MUSGS0137C2
AZ470253 1M0284E14
AZ853595 2M0156D23
AZ46466 1M0502C05
AZ966795 2M0156D23
AZ462250 T. brucei
AZ410166 1M018ZJ17
AZ710166 1M018ZJ17
AZ4453 AUZ54453
AZ659603 1M0537N05
AZ42423 1M0203C17
AZ796447 2M0051D09
AUU07217 AU007217
AU007217 AU007217
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/clone lib="Mouse 10kb plasmid UNGCIM library"
/note="Vector: PWD42rv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Biomedical Polymers Research Bldg., 20 S. 2030 E., SLC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0013 row: I column: 06
Seq primer: CGTTGTAAACGACGCCAGT
Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Laboratory Mouse DNA Resource
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Unpublished (2000)
Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /mol_type="genomic DNA"
/strain="C57BL/6J"
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/clone="UUGC1M0013106"
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Location/Qualifiers
                  AZ387817
AZ584023
AZ784029
AZ789794
AZ783417
D19150
AZ470253
AZ85395
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AZ309241.1 GI:10350030
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Tel: 801 585 5606
Fax: 801 585 7177
plasmid inserts
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84112, US
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VERSION
KEYWORDS
SOURCE
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AUTHORS
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SALK 1499
2M0065M14
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AZ428450 1M0210024
AZ368997 1M0119B20
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X88105 H.sapiens D
AZ445641 1M0241L05
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 T. brucei

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AG189412 I
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AU258068 AZ660125
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                  GenCore version 5.1.6
Copyright (c) 1993 - 2005 Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                                                                           34239544 segs, 19032134700 residues
                                                                                                                                                                                                                                                                                                                                                                                                                                  Total number of hits satisfying chosen parameters:
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Maximum Match 100%
Listing first 45 summaries
                                                                                                      nucleic search, using sw model
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Gapop 10.0 , Gapext 1.0
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gb_htc:;
gb_est3:;
gb_est4:;
gb_est6:;
gb_gs81:;
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Match Length
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DB seq length: 30
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Gaps

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Park, H., Kim, Y., Kim, S., Han, Y., Woo, T., Park, K., Eun, C.J.,
Hoon, S.T., Chu, M., Kim, H., Joo, S., Kim, C., Song, W. and Yoo, H.
Direct Submission
Submitted (07-JAN-2002) Hong-Seog Park, Korea Research Institute of
Bioscience and Biotechnology (KRIBB), Genome Research Center (GRC),
Eloscience and Biotechnology (KRIBB), Genome Research Center (GRC),
Eloscience and Biotechnology (KRIBB)
Eloscience and Biotechnology (KRIBB)
Eloscience and Biotechnology (KRIBB)
Eloscience and Biotechnology (KRIBB)
Cloner Research Center (GRC),
Tel:82-42-866-7181, Fax:82-42-860-4409)
Clones are derived from the chimpanzee BAC library RP-43 This BAC
end was generated during the R&D process and may have higher chance
of clone tracking errors.
                                                                                                                                                                                                                                         AG204232 27 bp DNA linear GSS 06-MAR-2004
Pan troglodytes DNA, clone: RP43-089J21.T7, genomic survey
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Pan troglodytes (chimpanzee)
Pan troglodytes
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
                                                                                                                                                                                                                                                                                                                                                                                                                             Pan troglodytes
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Pan.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Park, H., Kim, Y., Kim, S., Han, Y., Woo, T., Park, K., Eun, C.J., Hoon, S.T., Chu, M., Kim, H., Joo, S., Kim, C., Song, W. and Yoo, H. BAC end sequences of Library RP-43 Unpublished
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AG189412 20 bp DNA linear GSS 06-
Pan troglodytes DNA, clone: RP43-063018.TJ, genomic survey
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /cell_type="lymphocytes"
/clone_lib="RP-43 Chimpanzee Male BAC Library"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 27;
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/mol_type="genomic DNA"
/db_xref="taxon:9598"
             Mismatches
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R.Site 1 : ECORI
R.Site 2 : ECORI.
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                                                             1 CUGAGUUUAAAAGGCACC 18
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                                                                                                                CGGAGTACAAACGGCACC
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             13; Conservative
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AG189412
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was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymuclectide kinase. Adaptor oligonuclectides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWM92 (gi|4732114|gb|AR129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent B. coli XI10-GG1d (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CD532363 27-Dp mRNA linear EST 31-DEC-2003 26011 Arabidopsis Leaf Senescence Library Arabidopsis thaliana cDNA 3', mRNA sequence.
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/dev stage="Yellow Leaf With Greenish Base Area"
/dev stage="Yellow Leaf With Greenish Base Area"
/dab host="E. coli"
/clone lib="Arabidopsis Leaf Senescence Library"
/note="Organ: Rosette Leaf; Vector: pBluscript SKII+;
Site 1: EcoRK; Site 2: EcoRI; Senescent rosette leaves #5
and #6 (counted from the bottom) were harvested and
immediately frozen in liquid N2. The leaves were visibly
yellow excepted for the leaf base areas that were still
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Eukaryota, Viridiplantae, Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; rosids; eurosids II; Brassicales; Brassicaceae; Arabidopsis. I (bases I to 21).

Guo, Y., Cai, Z. and Gan, S.

Transcriptome of Arabidopsis leaf senescence
Plant Cell Environ. 27 (5), 521-549 (2004)
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119 Plant Science, Cornell University, Ithaca, NY 14853-5904, USA
Tel: 607 254 5418
Fax: 607 255 0599
Email: sg288@cornell.edu
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Pred. No. 2.1e+05;
2; Mismatches 3;
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|mol_type="manka"
|ecotype="Landsberg erecta"
|db_xref="taxon:3702"
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Insert Length: 25 Std Error: 0.00
Seq primer: T7
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Arabidopsis thaliana
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CD532363.1 GI:40452375
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KEYWORDS SOURCE ORGANISM

AUTHORS TITLE JOURNAL

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Query Match

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Laboratory Mouse DNA Resource
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                                                                                                                                         Park, H., Kim, Y., Kim, S., Han, Y., Woo, T., Park, K., Eun, C.J., Hoon, S.T., Chu, M., Kim, H., Joo, S., Kim, C., Song, W. and Yoo, H. Direct Submission

Direct Submission

Submitted (10-JAN-2002) Hong-Seog Park, Korea Research Institute of Submitted (10-JAN-2002) Hong-Seog Park, Korea Research Institute of Bioscience and Biotechnology (KRIBB), Genome Research Center (GRC); 52, Oun-dong, Yusong-gu, Daejeon 305-333, Korea

(E-mail:redstone@mail.kribb.re.kr, URL:http://phs.grc.kribb.re.kr/, Tel:82-42-866-7181, Fax:82-42-860-4409)

Clones are derived from the chimpanzee BAC library RP-43 This BAC end was generated during the R&D process and may have higher chance of clone tracking errors.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CC887301 30 bp DNA linear GSS 31-JUL-2003 SALK_149938.31.65.x Arabidopsis thaliana TDNA insertion lines Arabidopsis thaliana genomic clone SALK_149938.31.65.x, genomic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Alonso, J.M., Leisse, T.J., Barajas, P., Chen, H., Cheuk, R., Gadrinab, C., Jeske, A., Karnes, M., Kim, C.J., Parker, H., Prednis, L., Shinn, P., Zimmerman, J. and Ecker, J.R.
A Sequence-Indexed Library of Insertion Mutations in the
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This is single pass sequence recovered from the left border of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                      Park, H., Kim, Y., Kim, S., Han, Y., Woo, T., Park, K., Eun, C.J., Hoon, S.T., Chu, M., Kim, H., Joo, S., Kim, C., Song, W. and Yoo, H. BAC end sequences of Library RP-43
Mammalia; Butheria; Primates; Catarrhini; Hominidae; Pan.
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/clone_lib="RP-43 Chimpanzee Male BAC Library"
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Pred. No. 5.3e+05;
4; Mismatches 1; Indels
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Unpublished (2001)
Unpublished (2001)
Contact: Joseph R. Ecker
Salk Institute Genomic Analysis Laboratory (SIGnAL)
The Salk Institute for Biological Studies
Thoulo N. Torrey Pines Road, La Jolla, CA 92037, USA
Tel: 858 453 4100 x1752
Fax: 858 558 6379
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      organism="Pan troglodytes"
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Arabidopsis thaliana
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/db_xref="taxon:9598"
/clone="RP43-063018.TJ"
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/mol Lype="genomic DNA"
/ecctype="col-0"
/bo xref="taxon:3702"
/bo xref="taxon:3702"
/clone="SALK 19938.31.65.x"
/clone="SALK 19938.31.65.x"
/clone lib="Arabidopsis thaliana TDNA insertion lines ach of which contains one or more TDNA insertion elements. The resultant fragment for each line was directly sequenced to determine the genomic sequence at the site of insertion. Details of the protocols used cond at http://signal.salk.edu/tdna_protocols.html"
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Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhauseern, A. and Wright, D., Weiss, R.
Mouse whole genome scaffolding with paired end reads from 10kb plasmid inserts
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2M0065M14F Mouse 10kb plasmid UUGCIM library Mus musculus genomic clone UUGC2M0065M14 F, genomic survey sequence.
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Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Research Bldg., 20 S. 2030 E., SLC,
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/lab host="B. Coli strain XL10-Gold, T1-resistant, F-"
/clome lib="Mouse 10kb plasmid UUGCIM library"
/note="Vector: PWD42nr; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackson
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       53.3%; Score 11.2; DB 9; Length 30; 68.8%; Pred. No. 6.8e+05; ive 2; Mismatches 3; Indel8
                                     1. .30
/organism="Arabidopsis thaliana"
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Fax: 801 585 7177
Fax: 801 585 7177
Fax: 801 585 7177
Insert Length: 10000 Std Error: 0
Plate: 0065 row: M column: 14
Seq primer: CGTTGTAAAACGACGGCCAGT
Class: plasmid ends
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/strain="C57BL/6J"
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Location/Qualifiers
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/clone="UUGC2M0065M14"
Location/Qualifiers
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| CGGAGTATAAAAGGAA 17
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Best Local Similarity 68.8'
Matches 11; Conservative
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ORIGIN

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was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gil4732114 [gb] AR129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent B. coli XI10-GG1d (Stratagene) cells and selected for ampicillin resistance."
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2M0174J12F Mouse 10kb plasmid UUGC1M library Mus musculus genomic
clone UUGC2M0174J12 F, genomic survey sequence.
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Dunn, D., Aoyagi. A.)
Barber, M., Beacorn, T., Duval, B., Hamil, C., Reilly, M., Rose, R., Stokes, R., Menen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.
Mouse whole genome scaffolding with paired end reads from 10kb plasmid inserts
Unpublished (2000)
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/clone_lib="Mouse_10kb plasmid UDGCIM library"
/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackson
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University of Utah Genome Center
University of Utah
University of Utah
Rm. 308, Biomedical Polymers Research Bldg., 20 S. 2030 E., SLC,
84112, USA
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Insert Length: 10000 Std Error: 0.00
Plate: 0174 row: J column: 12
Seg primer: CGTTGTAAAACGACGGCCAGT
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/organism="Mus musculus"
/orlinel.commus DNA"
/strain="C57BL/6J"
/db xref="taxon:10090"
/clone="UUGC2M0174J12"
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Location/Qualifiers
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Fax: 801 585 7177
Email: ddunn@genet
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Best Local Similarity 6
...rhes 9; Conservat;
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AZ864535/c
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TITLE

COMMENT

FEATURES

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was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end repaired with T4 DNA polymerase and T4 polymuclectide kinase. Adaptor oligonuclectides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gi|4732114|gb|AR129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XI10-Gold (Stratagene) cells and selected for ampicillin resistance."
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Eukaryota; Metazoa; Chordata; Sciurognathi; Muridae; Murinae; Mus.
Eu (bases 1 to 25)
El (bases 1 to 25)
Ellan, H., Longacze, S., Mahmoud, M., Meenen, E., Pedersen, T.,
Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von
Niederhausern, A. and Wright, D., Weiss, R.
Mouse whole genome scaffolding with paired end reads from 10kb
plasmid inserts
Unpublished (2000)
Contact: Robert B. Weiss
University of Utah
University of Utah
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Insert Length: 10000 Std Error:
Plate: 0162 row: K column: 19
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Seq primer: CGTTGTAAAACGACGGCCAGT
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/strain="C57BL/6J"
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/clone="UUGC1M0162K19"
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High quality sequence stop: 25.
Location/Qualifiers
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Fax: 801 585 7177
Email: ddunn@genet
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AZ397442/c
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(http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gil 4712114 [gb]AP129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli Xil0-Gold (Stratagene) cells and selected for ampicillin resistance."
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Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Nelly, M., Rose, M., Rose, K., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.

Mouse whole genome scaffolding with paired end reads from 10kb
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/note="Vector: PWD42nv; Purified genomic DNA from M.musculus C57BL/6J (male) was obtained from the Jackson
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Length: 10000 Std Error: 0.00
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Seq primer: CACACAGGAAACAGCTATGACC
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Unpublished (2000)
Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
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/strain="C57BL/6J"
/db_xref="taxon:10090"
/clone="UUGC2M0043H11"
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High quality sequence stop: 25.
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Mus musculus
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24 TGGGTTTAAAAGAGAGC 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Tel: 801 585 5606
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Email: ddunn@gene
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(http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gelectrophoresis. Vector DNA was prepared from a derivative of pwaley (gql 4732114 [gb] API29072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XIIO-Gold (Stratagene) cells and selected for ampicillin resistance."
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Caston Cremieux, 91057 Evry cedex, FRANCE
PCR was performed on DNA from transformants of Arabidopsis thaliana plants from INRA (Versaillee). The DNA fragment(s) resulting from the PCR were directly sequenced from the left or the right border to determine the genomic sequence flanking the insertion. T-DNA derived sequences were removed. Information to order the corresponding mutant line and a link to a database providing a graphical display of the insertion site are available at http://dbsgap.versailles.inra.fr/publiclines/. This sequence has program 'Genoplante' (http://www.genoplante.com and http://genoplante-info.infoioingen.fr).
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Arabidopsis thaliana T-DNA flanking sequence, left border, clone
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AJS88176.1 GI:37937800
GSS; left border; T-DNA flanking sequence.
Arabidopsis thaliana (thale cress)
Arabidopsis thaliana
Arabidopsis thaliana
Bukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; rosids; eurosids II; Brassicales; Brassicaceae; Arabidopsis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Brunaud, V., Balzergue, S., Dubreucq, B., Aubourg, S., Samson, F., Chavvin, S., Bechtold, N., Cruaud, C., DeRose, R., Pelletier, G., Lepiniec, L., Caboche, M. and Lecharny, A.

T-DNA integration into the Arabidopsis genome depends on sequences
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/clone_lib="Arabidopsis thaliana T-DNA insertion lines"
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/cultivar="Wassillewskija"
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EMBO Rep. 3 (12), 1152-1157 (2002)
22363535
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VERSION KEYWORDS

SOURCE

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TITLE

FEATURES

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REFERENCE AUTHORS

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Mus musculus

Bukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,

Mamalia, Eutheria, Rodentia, Sciurognathi, Muridae, Murinae, Mus.

1 (bases 1 to 24)

Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C.,

Islam, H., Longacre, S., Mahmoud, M., Meenen, B., Pedersen, T.,

Reilly, M., Rose, M., Stokes, R., Tingey, A., von

Niederhausern, A. and Wright, D., Weiss, R.,

Mouse whole genome scaffolding with paired end reads from 10kb

plasmid inserts
                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Mammalia; Butheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
1 (bases 1 to 24)
Kato,K. and Matoba,R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AZ660125
1M0538F04F Mouse 10kb plasmid UUGCIM library Mus musculus genomic clone UUGCIM0538F04 F, genomic survey sequence.
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                                                                                                                                      Generation of expressed sequence tags from mouse brain Unpublished (2002)
Contact: Kikuya Kao.
Contact: Kikuya Kao.
Graduate School of Biological Sciences
Nara Institute of Science and Technology
Self-5 Takayama, Ikoma, Nara 630-0101, Japan
Tel: 81-743-72-5581
Eax: 81-743-72-5589
Email: Kkato@bs.aist-nara.ac.jp,
UKL:http://lvow2.aist-nara.ac.jp,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   49.5%; Score 10.4; DB 1; Length 24; 66.7%; Pred. No. 1.7e+06; ive 3; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /mol_type="mRNA"
/db_xref="taxon:10090"
/clone="BED0011993"
/tissue_type="brain"
/clone_lib="3'-directed mouse cDNA library"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Tel: 801 585 5606
Fax: 801 585 7177
Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00

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    /organism="Mus musculus"

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Class: plasmid ends
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Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
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/strain="C57BL/6J"
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  musculus (house mouse)
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If (bases 1 to 23)

Hall,N. J. Bowman,S., Lennard,N.J., Doggett,J., Atkin,R.,

Chillingworth,C., Ormond,D., Harris,B., El-Sayed,N., Hou,L.,

Chillingworth,C., Ormond,D., Harris,B., El-Sayed,N., Hou,L.,

Melville,S.E., Rajandream,M.A. and Barrell,B.G.

Direct Submission

Submitted (10-DEC-2000) Trypanosoma brucei genome sequencing

project, Sanger Centre, The Wellcome Trust Genome Campus, Hinxton,

Cambridge CB10 1SA, E-mail: barrell@sanger.ac.uk and

nhl@sanger.ac.uk

Constructed at the Institute for Genomic Research (TIGR),

Rockville, MD. Genomic DNA isolated from a cloned population of

Trypanosoma brucei (TREU927/4 GUTAt 10.1) was mechanically sheared

to give a tight size distribution (

4 kb). The v + i method used for the library construction is

described in detail in Smith, H. and Venter, J.C. (Making small

insert libraries for whole genome shotgun sequencing projects.

Barrell, Oxford University Press, 1999).
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AU258068 1 GI:20323293
                                                                                                                                                                                                                                                                                                                                                                                           TA130G11Q 23 bp DNA linear GSS 13-DEC-2000 T. brucei sheared genomic DNA clone 130g11, reverse sequence, genomic survey sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Email: nelsayed@tigr.org
Details of T. brucei sequencing at the Sanger Centre are available
at http://www.sanger.ac.uk/Projects/T_brucei/.
Location/Qualifiers
                                                                                                                                                                          Gaps
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Eukaryota, Euglenozoa, Kinetoplastida, Trypanosomatidae,
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                                                                                                                       Length 28;
                                                                                                                                                                          4; Indels
                                                                                                                       Query Match

So.5%; Score 10.6; DB 9;
Best Local Similarity 58.8%; Pred. No. 1.4e+06;
Matches 10; Conservative 3; Mismatches 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /organism="Trypanosoma brucei"
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/strain="TREU927"

    .28
/note="T-DNA flanking sequence
left border"

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                                                                                                                                                                                                                        1 CUGAGUUUAAAAGGCAC 17
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Best Local Similarity 66.7
Matches 8; Conservative
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  misc_feature
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TA130G11Q/c
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84112, USA
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Bukaryota; Metazoa; Chordata; Sciurognathi; Muridae; Murinae; Mus.

Talam, Butheria; Rodentia; Sciurognathi; Muridae; Mus.

Bunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C.,

Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T.,

Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von

Niederhausern, A. and Wright, D., Weiss, R.

Mouse whole genome scaffolding with paired end reads from 10kb

plasmid inserts

Unpublished (2000)

Contact: Robert B. Weiss

University of Utah Genome Center

University of Utah

Rm. 308, Biamedical Polymers Research Bldg., 20 S. 2030 E., SLC, UT

##112, Aug. Brand.
                                                                                                              (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligomucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gal electrophoresis. Vector DNA was prepared from a derivative of pwDAQ2 (gql 4732114[gbl A722072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XII.0-Gold (Stratagene) cells and selected for ampicillin resistance."
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Insert Length: 10000 Std Error: 0.00
Plate: 0261 row: C column: 20
Seq primer: cGTTGTAAAACGACGCCAGT
Class: plasmid ends
                                                                                                  Laboratory Mouse DNA Resource

    .29
    /organism="Mus musculus"

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/strain="C57BL/6J"
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Best Local Similarity 58.3°
7; Conservative
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Fax: 801 585 7177
Email: ddunn@genet
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22 TGTGTTTAAAG 11
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muscular C7) Muse DNR Resource (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared bunk was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarese gel electrophoresis. Vector DNA was prepared from a derivative of pwD42 (gi|4732114|gb|AF129072.1) a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli X110-GG1d (Stratagene) cells and selected for ampicillin resistance."
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Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacrer, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.
Mouse whole genome scaffolding with paired end reads from 10kb plasmid inserts
Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AZ813693 10Kb plasmid UUGCIM library Mus musculus genomic clone UUGC2M0081P03F Pt. genomic survey sequence.
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/clone_11b="Mouse lOkb plasmid uUGCNM library"
/note="Vector: PWD47nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackson
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Fax: 801 585 7177
Email: ddunn@genetics.utah.edu
Insert Length: 1000 Std Error: 0
Plate: 0081 row: P column: 03
Seg primer: CGTTGTAAAACGACGGCCAGT
Class: plasmid ends
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/strain="C57BL/6J"
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Location/Qualifiers
xref="taxon:10090'
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20 TCAGTTTAAAAG 9
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/lab.host="B" Coli strain XL10-Gold, T1-resistant, F-"
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/note="Vector: PWD42n; Purified genomic DNA from M.
musculus G57BL/6/ (male) was obtained from the Jackson
Laboratory Mouse DNA Resources (documents/dnares/). The DNA
was bydrodynamically sheared by repeated passage through a
0.005 inch orifice at constant velocity. The sheared DNA
was hydrodynamically sheared by repeated baseage through a
0.005 inch orifice at constant velocity. The sheared DNA
was blunt end-repaired with T4 DNA polymerase and T4
polymuclectide kinase. Adaptor oligonuclectides were
ligated to the blunt ends in high molar excess. The
adaptored DNA was purified and size-selected for a 9.5 to
10.5 kb range using preparative agarose gel
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of pWD42 (gil 4732114|gp| AR129072.1); a copy-number
inducible derivative of plasmid R1. The vector was ligated
with adaptors complementary to the insert adaptors and
purified. The sheared, adaptored mouse DNA was annealed to
adaptored vector DNA, and transformed into
chemically-competent E. coli Xillo-Gold (Stratagene) cells
and selected for ampicillin resistance."
xref="taxon:10090"
                                           /clone="UUGC2M0081P03"
/sex="Male"
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ORIGIN

Gaps , 0 49.5%; Score 10.4; DB 8; Length 29; 58.3%; Pred. No. 1.7e+06; tive 4; Mismatches 1; Indels Query Match
Best Local Similarity 58.3
Matches 7; Conservative

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1 CUGAGUUUAAAA 12 |:|| :::|||| 12 CTGACTTTAAAA 23 12

ò g Search completed: August 19, 2005, 01:37:26 Job time : 2980 secs

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August 18, 2005, 22:25:38 ; Search time 416 Seconds (without alignments) 298.833 Million cell updates/sec
GenCore version 5.1.6
Copyright (c) 1993 - 2005 Compugen Ltd.
                                                                                            - nucleic search, using sw model
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1 cugaguuuaaaaggcacccnn 21 US-10-665-951-2185 21 Perfect score: Sequence: Title:

IDENTITY_NUC Gapop 10.0 , Gapext 1.0 Scoring table:

4390206 seqs, 2959870667 residues Searched:

Total number of hits satisfying chosen parameters:

3522762

Minimum DB seq length: 0 Maximum DB seq length: 30

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

geneseqn2000s:* geneseqn2001as:* geneseqn2001bs:* geneseqn2002as:* geneseqn2003cs:* genesegn2003ds:* geneseqn2003as:* geneseqn2003bs:* N_Geneseq_16Dec04:* geneseqn2002bs:* geneseqn2004as: geneseqn1990s:* genesegn1980s: •• Database

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

geneseqn2004bs:

SUMMARIES

Result No. No. Sc. 2 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	8000 119 119 119 119 119 119 119 119	A C C C C C C C C C C C C C C C C C C C	Match Length DB Match Length DB 90.5 5 21 9 90.5 5 21 9 90.5 5 21 9 90.5 5 21 19 90	10000000000000000000000000000000000000	ID ADA13968 ADA13964 ADA13613 ADA13613 ADE37703 ADE37751 ADE37775 ADE37775 ADE377907 ADE3890	Description Ada1366 Short int Ada1364 Short int Ada13613 Short int Ada1360 Short int Adf37751 Human VEG Adf37751 Human VEG Adf3775 Human VEG Adf3775 Human VEG Adf3776 Human VEG Adf37789 Human VEG Adf37799 Fuman VEG Adf3779 Fuman VEG
118 20	11 1 1 1 1 1 1 1 1 1	90.08	217	2222	ADG29896 ADG29828 ADG29881	

		Adf37772 Human VEG	Adf37812 Human VEG	Adf37742 Human VEG		Adg29490 FLT1 BiNA	Adg29893 FLT1-targ	Adg29917 FLT1-targ	Adg29887 FLT1-targ	fouse	Aax68744 Human flt	Acc86729 Human VEG	Aax68743 Human flt	Aax74492 Mouse flt	Acc86730 Human VEG	Aax68742 Human flt	. Aax74491 Mouse flt	Aaz87380 Human thr	Aaz87381 Human thr	Aax28169 PCR prime	Aat58464 PEPCK gen	Aav82574 PCR prime	Adf35731 Human VEG	Adf36158 Human VEG
ADG29920	ADG29824	ADF37772	ADF37812	ADF37742	ADF37748	ADG29490	ADG29893	ADG29917	ADG29887	AAX74493	AAX68744	ACC86729	AAX68743	AAX74492	ACC86730	AAX68742	AAX74491	AAZ87380	AAZ87381	AAX28169	AAT58464	AAV82574	ADF35731	ADF36158
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21	21	23	23	23	23	23	23	23	23	17	17	20	17	17	20	17	17	22	24	28	30	30	19	19
90.5	90.5	90.5	90.5	90.5	90.5	90.5	90.5	90.5	90.5	81.0	81.0	81.0	76.2	76.2	76.2	71.4	71.4	67.6	9.19	65.7	62.9	62.9	61.9	61.9
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ALIGNMENTS

RESULT 1

short interfering nucleic acid; šiNA; expression; replication; inhibition; RNA interference; virucide; anti-HTV; hepatorropic; anti-HTV; hepatorropic; autiinflammatory; plant; anti-Yral; vasotropic; neuroprotective; cytostatic; cardiovascular; immunosuppressive; respiratory; nephrotropic; endocrine; viral infection; hepatitis B; hepatitis C; HTV; herpes simplex; cytomegalovirus; hepatitis C; HTV; respiratory syncytial virus; influenza virus; restenosis; neurodegeneration; cancer; neurological; prion; inflammatory; autoimmune; pulmonary; renal; liver; mitochondrial; reproductive disease; Short interfering nucleic acid (siNA) oligonucleotide SEQ ID NO:305. double-stranded short interfering nucleic acid; BP. ADA13968/c ID ADA13968 standard; RNA; 21 (first entry) pulmonary; renal; liver; m chemical modification; ss. 20-NOV-2003 Synthetic. ADA13968;

WO2003070918-A2.

20-FEB-2003; 2003WO-US005346 28-AUG-2003

2002US-0363124P. 2002US-0386782P. 2002US-0406784P. 2002US-0408378P. 09-SEP-2002; 2002US-0409293P. 2002US-0358580P 06-JUN-2002; 29-AUG-2002; 05-SEP-2002; 09-SEP-2002; 20-FEB-2002; 11-MAR-2002;

(RIBO-) RIBOZYME PHARM INC

Pavco P; Beigelman L, Macejak D, Zinnen S, Fosnaugh K, Mokler V, Jamison S; Mcswiggen J, Morrissey D,

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the sink molecule comprises a consider solution in the constraint dewrited as a consider solution of a target gene, where the sink molecule comprises about 21 nucleotides. Also described: (1) a sink molecule that inhibits expression of target RNA; (2) a sink molecule that inhibits expression of atarget RNA; (2) a sink molecule that inhibits expression of a virus and optionally does not require presence of a ribonucleotide for inhibition; (3) a sink molecule that inhibits expression of a target gene and does not require presence of a ribonucleotide for inhibition; (4) a sink molecule that inhibits expression of a target gene and does not require presence of a ribonucleotide for inhibition; (4) a sink molecule that inhibits molecules. Sink's can have virucide, anti-HIV, hepatotropic, anti-HIM and anti-HIV, hepatotropic, anti-HIM and anti-HIV, hepatotropic, anti-HIM and anti-HIV, hepatotropic, anti-HIM and activities. The sink's are useful for downsegulating and endocrine activities. The sink's are useful for downsegulating and endocrine activities. The sink's are useful for downsegulating cylcatic, cardiovagoular, immunosuppression of target RNA, and inhibiting replication of a virus. Sink molecules can be used: (a) for therapy of any disorder that responds to modulation of gene expression, especifically and plant viral infections, specifically and plant viral infections.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      C; HIV; herpes simplex; cycomegalo; human papilloma; respiratory syncytial or influenza viruses, and also many other diseases such as restenosis, neurodegeneration, cancers, and cardiovascular, neurological, prion, inflammatory, autoimmune, pulmonary, renal, liver, mitochondrial, endocrine or reproductive diseases; and (b) for diagnosis, target avalidation, genomic discovery, genetic engineering, pharmacogenomics and analysis of gene function. Chemical modification of siNA molecules improves interfering activity; stability; cellular uptake; binding affinity and/or mediates increased polymerase activity. siNA may be designed to target many related genes containing a conserved sequence. The present sequence represents a siNA oligomucleotide sequence, which is used in the exemplification of the present invention.
                                               New short interfering nucleic acid containing no ribonucleotides, useful
e.g. for treating viral infection, downregulates expression of target
gene or RNA.
                                                                                                                                                                                                                                                                               present invention describes a double-stranded short interfering
                                                                                                                                                                                                                 Example 4; Page 142; 204pp; English.
WPI; 2003-689785/65
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Sequence 21 BP; 4 A; 4 C; 5 G; 2 T; 6 U; 0 Other;

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90.5%; Score 19; DB 9; Length 21; 78.9%; Pred. No. 1.8; 1.ve 4; Mismatches 0; Indels
                                                 CUGAGUUUAAAAGGCACCC 19
                                                               15; Conservative
            Similarity
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 Query Match
Best Local 8
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0; Gaps

RESULT 2

Short interfering nucleic acid (siNA) oligonucleotide SEQ ID NO:301.

short interfering nucleic acid; siNA; expression; replication; hinbition; RNA interference, virucide; anti-HIV; hepatotropic; anti-inflammatory; plant; anti-inviral; vasotropic; neuroprotective; cytostatic; cardiovascular; immunosuppressive; respiratory; nephrotropic; endocrine; viral infection; hepatitis B; hepatitis C; HIV; herpes simplex; cytomegalovirus; human papillomavirus; respiratory syncytial virus; influenza virus; restenosis; neurodegeneration; cancer; neurological; prion; inflammatory; autoimmune; pulmonary; renal; liver; mitochondrial; reproductive disease; chemical modification; ss. double-stranded short interfering nucleic acid;

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the present invention describes a double-stranded short interfering nucleic acid (siNA) that downregulates expression of a target gene, where the siNA molecule comprises no ribonuclectides and each strand of the double-stranded siNA comprises about 21 nuclectides. Also described: (1) a siNA molecule that inhibits expression of a virus and optionally does not require presence of a ribonuclectide for inhibition; (3) a siNA molecule that inhibits replication of a virus and optionally does not require presence of a ribonuclectide for inhibition; (4) a siNA molecule that inhibits expression of a target gene and does not require presence of a ribonuclectide for inhibition; (4) a siNA molecule that inhibits expression of a target gene by mediating RNA interference; and (5) a molecule siNA's can have virucide, anti-HIV hepatchropic, antiinflammatory, plant antiviral, vasotropic, neuroprotective, cytostatic, cardiovascular; immunosuppressive, respiratory, nephrotropic, and endocrine activities. The siNA's are useful for downregulating expression of target genes, inhibiting replication of a virus. SiNA molecules can be used (a) for therapy of any disorder that responds to modulation of gene expression of target simplex; cytomegalo; human papilloma; respiratory especially animal and plant viral infections, specifically hepatitis of cytomegalo; human papilloma; respiratory cytial or influenza viruses, and cardiovascular, neurological, prior, inflammatory, autoimmune, pulmonary, renal, liver, mitochondrial, cendocrine or reproductive diseases; and cardiovascular, neurological, prior, inflammatory, autoimmune, pulmonary, renal, liver, mitochondrial, genomic discovery, genetic engineering pharmacogenemics and analysis of gene function. Chemical modification of siNA molecules can engerent sequence represente asiNA chemical polymerase activity; stability; cendocrine or repro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New short interfering nucleic acid containing no ribonucleotides, useful
e.g. for treating viral infection, downregulates expression of target
gene or RNA.
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Morrissey D, Fosnaugh K, Mokler V, Jamison S;
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Best Local Similarity 78.3.
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                                         WO2003070918-A2
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Synthetic.
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                                                                                                                                              short interfering nucleic acid; siNA; expression; replication; inhibition; RNA interference; virucide; anti-HV; hepatorropic; antiinflammatory; plant; antiviral; vasotropic; neuroprotective; cytostatic; cardiovascular; immunosuppressive; respiratory; nephrotropic; endocrine; viral infection; hepatitis B; hepatitis C; HIV; herpes simplex; cytomegalovirus; human papillomavirus; respiratory syncytial virus; influenca virus; restenosis; neurodegeneration; cancer; neurological; prion; inflammatory; autoimmune; pulmonary; renal; liver; mitochondrial; reproductive disease;
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prion, inflammatory, autoimmune, pulmonary, renal, liver, mitochondrial,
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                                                                                                     Short interfering nucleic acid (siNA) oligonucleotide SEQ ID NO:345.
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Mokler V, Jamison S;
                                                                                                                                  double-stranded short interfering nucleic acid;
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2002US-0406784P.
2002US-0408378P.
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Fosnaugh K,
              ADA13613 standard; RNA; 21
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                                                                         (first entry)
                                                                                                                                                                                                                                                                                     chemical modification; ss
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06-JUN-2002; 2
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05-SEP-2002; 2
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15-JAN-2003;
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Morrissey D,
                                                                         20-NOV-2003
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                                           ADA13613
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ADA13613
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                    validation, genemic discovery, genetic engineering, pharmacogenomics and analysis of gene function. Chemical modification of sinA molecules improves interfering activity; stability; cellular uptake; binding affinity and/or mediates increased polymerase activity. sinA may be designed to target many related genes containing a conserved sequence. The present sequence represents a sinA oligonucleotide sequence, which is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   short interfering nucleic acid; siNA; expression; replication; inhibition; RNA interference; virucide; anti-HIV; hepatotropic; anti-HIV; hepatotropic; anti-HIV; neuroprotective; cytostatic; cardiovacular; immunosuppressive; respiratory; nephrotropic; endocrine; viral infection; hepatitis B; hepatitis C; HIV; herpes simplex; cytomegalovirus; human papillomavirus; respiratory syncytial virus; influenza virus; restenosis; neurodegeneration; dancer; neurological; prion; inflammatory; autoimmune; pulmonary; renel; liver; mitochondrial; reproductive disease;
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reproductive diseases; and (b) for diagnosis, target
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Fosnaugh K, Mokler V, Jamison S;
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                                                                                                                                                                                                                                                                                                        DB 9;
                                                                                                                                                                                                                                                                                                      Score 19; DB 9
Pred. No. 1.8;
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2003US-0440129P.
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  endocrine or
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Morrissey D,
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05-SEP-2002;
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double-stranded sinA comprises about 21 nucleotides. Also described: a sinA molecule that inhibits expression of target RNA; (2) a sinA molecule that inhibits expression of target RNA; (2) a sinA molecule that inhibits expression of a virus and optionally does not require presence of a ribonucleotide for inhibition; (4) a sinA molecule that inhibits expression of a target gene and does not require presence of a ribonucleotide for inhibition; (4) a sinA molecule that inhibits expression of a target gene by mediating RNA interference; and (5) a method for modulating expression of a gene in a cell using sinA molecules. SinA's can have virucide, anti-HIV, hepacotropic, anti-HIM manatory, plant anti-viral, vasotropic, neuroprotective, cytostatic, cardiovascular, immunosuppressive, respiratory, nephrotropic and endocrine activities. The sinA's are useful for downregularing carpression of target RNA, and inhibiting replication of a virus. SiNA molecules can be used: (a) for therapy of any disorder that responds to modulation of gene expression, therapy of any disorder that responds to modulation of gene expression, especially animal and plant viral infections, specifically hepatitis B or synchrial or influenza viruses, and also many other diseases such as
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    prion, inflammatory, autoimmune, pulmonary, renal, liver, mitochondrial, endocrine or reproductive diseases; and (b) for diagnosis, target validation, genomic discovery, genetic engineering, pharmacogenomics and analysis of gene function. Chemical modification of siNA molecules improves interfering activity, stability; cellular uptake; binding affinity and/or mediates increased polymerase activity. siNA may be designed to target many related genes contening a conserved sequence. The present sequence represents a siNA oligonucleotide sequence, which is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            double-stranded short interfering nucleic acid; short interfering nucleic acid; siNA; downregulation; sacular acid; siNA; downregulation; constant endothelial growth factor receptor; VEGFR; antiangiogenic; cytostatic; antidiabetic; ophthalmological; antiarthritic; antipsoriatic; nephrotropic; gynaecological; angiogenesis-associated condition; cancer; diabetic retinopathy; macular degeneration; neovascular glaucoma; arthritis; psoriasis; endometriosis; angiofibroma; polycystic kidney disease; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                        restenosis, neurodegeneration, cancers, and cardiovascular, neurological,
siNA molecule comprises no ribonucleotides and each strand of the
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Pred. No. 1.8;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             in the exemplification of the present invention.
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ive 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       90.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 19; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO2003070910-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               used
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20-FEB-2002; 2002US-0358580P. 11-MAK-2002; 2002US-0363124P. 29-MAX-2002; 2002WO-US017674. 6-JUN-2002; 2002US-0386782P.

20-FEB-2003; 2003WO-US005022

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The present invention describes a double-stranded short interfering nucleic acid (siNA) that downregulates expression of the vascular endothelial growth factor receptor (VEGFR) gene. Also described: (1) a siNA that downregulates the VEGF gene, (2) kits for in vitro or in vivo delivery of siNA; (3) conjugates and/or complexes of siNA; (4) vectors that express siNA, and (5) single-stranded siNA with similar properties. The siNAs have antiangiogenic, cytostatic, antidiabetic, ophthalmological, antiarthritic, antipsoriatic, nephrotropic and synascological activities. The siNA are useful for modulating (downregulating) the expression of VEGFR genes. The siNA are potentially useful for treating a wide range of angiogenesis-associated conditions, particularly cancers, diabetic retinopathy, macular degeneration, neovascular glaucoma, arthritis, psoriasis, endometriosis, anglofibroma, and polycystic kidney disease. The siNA may also be useful for diagnosis, diapeting, studying gene function, and also for gene mapping (e.g. of single-incleotide polymorphisms). The present sequence is used in the
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                                                                                                                                                                                                                                                                                       New double-stranded interfering nucleic acid, useful e.g. for treatment and diagnosis of cancer, downregulates the vascular endothelial growth factor receptor gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 21 BP; 6 A; 5 C; 4 G; 2 T; 4 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                       Example 3; SEQ ID NO 2100; 207pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              exemplification of the present invention.
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            29-JJJ-2002; 2002US-0399348P.
29-AUG-2002; 2002US-0406784P.
05-SEP-2002; 2002US-0409293P.
04-NOV-2002; 2002US-0028949.
27-NOV-2002; 2002US-00306747.
15-JAN-2003; 2003US-0440129P.
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                                                                                                                                                                         (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                Mcswiggen J, Beigelman L,
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ID ADF37751 standard; RNA; 21
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Best Local Similarity 100.
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The present invention describes a double-stranded short interfering nucleic acid (siNA) that downrequlates expression of the vascular endothelial growth factor receptor (VEGFR) gene. Also described: (1) a siNA that downrequlates the VEGF gene, (2) kits for in vitro or in vivo delivery of siNA; (3) conjugates and/or complexes of siNA; (4) vectors that express siNA; and (5) single-stranded siNA; (4) vectors of that express siNA; and (5) single-stranded siNA; with similar properties. The siNAs have antianglogenic, cytostatic, antidiabetic, ophthalmological, antiarthritic, antipsoriatic, nephrotropic and synamological activities. The siNA sere useful for modulating (downregulating) the expression of VEGFR genes. The siNA are potentially useful for treating a wide range of angiogenesis-associated conditions, particularly cancers, diabetic retinopathy, macular degeneration, neovascular glaucoma, arthritis, psoriasis, endometriosis, angiofibroma, and polycystic kidney disease. The siNA may also be useful for diagnosis, and polycystic kidney disease. The siNA may also be useful for diagnosis, engineering, studying gene function, and also for gene mapping (e.g. of single-nucleotide polymorphism). The present sequence is used in the
              cytostatic, antidiabetic; ophthalmological; antiarthritic; antiphoriatic; nephrotropic; gynaecological; angiogenesis-associated condition; cancer; diabetic retinopathy; macular degeneration; neovascular glaucoma; arthritis; psoriasis; endometriosis; angiofibroma; polycystic kidney disease; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New double-stranded interfering nucleic acid, useful e.g. for treatment and diagnosis of cancer, downregulates the vascular endothelial growth
endothelial growth factor receptor; VEGFR; antianglogenic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        90.5%; Score 19; DB 10; Length 21; 78.9%; Pred. No. 1.8;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mcswiggen J, Beigelman L, Pavco P;
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2002US-0386782P.
2002US-0393796P.
2002US-0399348P.
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2002US-0408378P.
2002US-0409293P.
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                                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                         29-MAY-2002;
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                                                                                                                                                                  Synthetic
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ADF37775/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention describes a double-stranded short interfering nucleic acid (siNA) that downregulates expression of the vascular endothelial growth factor receptor (VEGFR) gene. Also described: (1) a siNA that downregulates the VEGF gene. Also described: (1) a call which that downregulates the VEGF gene; (2) kits for in vitro or in vivo delivery of siNA; (3) conjugates and/or complexes of siNA; (4) vectors that siNAs have antianglogenic, cytostatic, antidiabetic, ophthalmological, antiarthritic, antipsoriatic, nephrotropic and cytostatic, antidiabetic, ophthalmological activities. The siNA are useful for modulating (downregulating) the expression of VEGFR genes. The siNA are potentially useful for treating a wide range of angiogenesis-associated conditions, particularly cancers, diabetic retinopathy, macular degeneration, and polycystic kidney disease. The siNA may also be useful for diagnosis, and polycystic kidney disease. The siNA may also be useful for diagnosis, drug screening, target identification and validation, genetic cangineering, studying gene function, and also for gene mapping (e.g. of single-nucleotide polymorphisms). The present sequence is used in the exemplification of the present invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      double-stranded interfering nucleic acid, useful e.g. for treatment diagnosis of cancer, downregulates the vascular endothelial growth
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short interfering nucleic acid; siNA; downregulation;
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ADP37745/C
ID ADP37745 standard; RNA; 21 BP.
XX
AC ADF37745;
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DT 12-FEB-2004 (first entry)
XX
DE Human VEGFR1 short interfering n
XX
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KW
Gouble-stranded short interfering n
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KW
Ghort interfering nucleic acid;
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2002US-0363124P.
2002WO-US017674.
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05-SEP-2002; 2002US-0408378P.
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15-JAN-2003; 2003US-0440129P.
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Best Local Similarity 78.5
مادات 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         factor receptor gene,
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    WO2003070910-A2
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29-JUL-2002;
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29-MAY-2002;
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CUGAGUUUAAAAGGCACCC

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CTGAGTTTAAAAGGCACCC

BP

ADF37899 standard; RNA; 21

1.8; Mismatches

78.98;

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double-stranded short interfering nucleic acid; short interfering nucleic acid; siNA; downregulation; sebort interfering nucleic acid; siNA; downregulation; vascular endothelial growth factor receptor; VEGFR; antiangiogenic; cytostatic; antidiabetic; ophthalmological; antiarchritic; antipsoriatic; nephrotropic; gynaecological; angiogenesis-associated condition; cancer; diabetic retinopsthy; meanular degeneration; neovascular glaucoma; arthritis; psoriasis; endometriosis; angiofibroma;
Best Local Similarity 78.9
Matches 15, Conservative
                                                                                            ADF37899/c
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                                                                                                                short interfering nucleic acid; siNA; downregulation; vascular endothelial growth factor receptor; VEGFR; antiangiogenic; cytostatic; antidiabetic; ophthalmological; antiarthritic; antipsoriatic; nephrotropic; gynaecological; angiogenesis-associated condition; cancer; diabetic retinopathy; macular degeneration; neovascular glaucoma; arthritis; psoriasis; endometriosis; angiofibroma; polycystic kidney disease; ss.
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                                                            Human VEGFR1 short interfering nucleic acid (siNA) SEQ ID NO:2188.
                                                                                double-stranded short interfering nucleic acid;
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    ВР
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09-SEP-2002; 2002US-0409293P.
04-NOV-2002; 2002US-00287949.
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2002US-0363124P.
2002WO-US017674.
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03-JUL-2002; 2002US-0393796P.
29-JUL-2002; 2002US-0399348P.
29-AUG-2002; 2002US-0406784P.
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    21
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                                          (first entry)
    ADF37775 standard; RNA;
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29-MAY-2002;
                                                                                                                                                                                 Homo sapiens
                                          12-FEB-2004
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                                                                                                                                                                        Synthetic
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Human VEGFR1 short interfering nucleic acid (siNA) SEQ ID NO:2096.

(first entry)

12-FEB-2004

ADF37899;

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nucleic acid (siNA) that downregulates expression of the vaccular endochelial growth factor receptor (VEGFR) gene. Also described: (1) a siNA that downregulates the VEGF gene; (2) kits for in vitro or in vivo delivery of siNA; (3) conjugates and/or complexes of siNA; (4) vectors that express siNA; and (5) single-stranded siNA with similar properties. The siNAs have antiangiogenic, cytostatic, antidiabetic, ophthalmological, antiathritic, antipsoriatic, nephrotropic and gynaecological activities. The siNA are useful for modulating (dwnregulating) the expression of VEGFR genes. The siNA are potentially useful for treating a wide range of angiogenesis-associated conditions, particularly cancers, diabetic retinopathy, macular degeneration,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          present invention describes a double-stranded short interfering
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2002US-0393796P.
2002US-0406784P.
2002US-0408378P.
2002US-0408378P.
2002US-0409293P.
2002US-003067449.
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nucleic acid (siNn). The control account of the vascular candothelial growth factor receptor (VEGFR) gene. Also described: (1) a siNA that downregulates the VEGF gene; (2) kits for in vitro or in vivo delivery of siNA; (3) conjugates and/or complexes of siNA; (4) vectors that express siNA; and (5) single-stranded siNA with similar properties. The siNAs have antiangiogenic, cytostatic, antidiabetic, ophthalmological, antiatrhritic, antiploriatic, nephrotropic and gynaecological, antiatrhritic, antiploriatic, nephrotropic and gynaecological activities. The siNA are useful for modulating (downregulating) the expression of VEGFR genes. The siNA are potentially useful for treating a wide range of angiogenesis-associated conditions, particularly cancers, diabetic retinopathy, macular degeneration, particularly cancers, diabetic retinopathy, macular degeneration, and polycystic kidney disease. The siNA may also be useful for diagnosis, and polycystic kidney disease. The siNA may also be useful for diagnosis, consisting and polycystic kidney disease. The siNA may also be useful for diagnosis, and polycystic kidney disease. The siNA may also be useful for diagnosis, and polycystic kidney disease. The siNA may also be useful for diagnosis, and polycystic kidney disease. The siNA may also be useful for diagnosis, and polycystic kidney disease. The siNA may also be useful for diagnosis, and polycystic kidney disease. The siNA may also be useful for diagnosis, and polycystic kidney disease. The siNA may also be useful for diagnosis, and polycystic kidney disease. The siNA may also be useful for diagnosis, and polycystic kidney disease. The siNA may also be useful for diagnosis, and polycystic kidney disease. The siNA may also be useful for diagnosis, and polycystic kidney disease. The siNA may also be useful for diagnosis, and polycystic kidney disease.

present invention describes a double-stranded short interfering

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(e.g. o in the

engineering, studying gene function, and also for gene mapping single-nucleotide polymorphisms). The present sequence is used exemplification of the present invention.

Sequence 21 BP; 4 A; 4 C; 5 G; 2 T; 6 U; 0 Other;

90.5%; Score 19; DB 10; Length 21;

Query Match

2002WO-US017674. 2002WO-US017674. 2002US-0386782P.

06-JUN-2002;

20-FEB-2003; 2003WO-US005022

20-FEB-2002;

WO2003070910-A2 Homo sapiens.

Synthetic.

28-AUG-2003

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                                                                                                                                                                                                                                                                                                                                                                                                                      double-stranded short interfering nucleic acid; short interfering nucleic acid; siNA; downregulation; selecting nucleic acid; siNA; downregulation; antiangiogenic; vascular endothelial growth factor receptor; VEGFR; antiangiogenic; cytostatic; antidiabetic; ophthalmological; antiathritic; antipsoriatic; nephrotropic; gynaecological; angiogenesis-associated condition; cancer; diabetic retinopsthy; macular degeneration; neovascular glaucoma; arthritis; psoriasis; endometriosis; angiofibroma;
neovascular glaucoma, arthritis, psoriasis, endometriosis, angiofibroma, and polycystic kidney disease. The siNA may also be useful for diagnosis, drug screening, target identification and validation, genetic engineering, studying gene function, and also for gene mapping (e.g. of single-nucleotide polymorphisms). The present sequence is used in the exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New double-stranded interfering nucleic acid, useful e.g. for treatment and diagnosis of cancer, downregulates the vascular endothelial growth
                                                                                                                                                                   Gaps
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                                                                                                                                                                                                                                                                                                                                                                                             Human VEGFR1 short interfering nucleic acid (siNA) SEQ ID NO:2104.
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                                                                                                                                  Score 19; DB 10; Length 21;
Pred. No. 1.8;
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                                                                                                        Sequence 21 BP; 4 A; 4 C; 5 G; 2 T; 6 U; 0 Other;
                                                                                                                                                                  4; Mismatches
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Best Local Similarity 78.99,
Best Local 15; Conservative
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06-JUN-2002;
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cendothelial growth factor receptor (VEGPR) gene. Also described: (1) a sinA that downregulates the VEGF gene; (2) kits for in vitro or in vivo delivery of sinA; (3) conjugates and/or complexes of sinA; (4) vectors that express sinA; and (5) single-stranded sinA with similar properties. The sinAs have antianthritic, antipsoriatic, nephrotropic and gynaecological, antiarthritic, antipsoriatic, nephrotropic and gynaecological activities. The sinA are useful for modulating (downregulating) the expression of VEGFR genes. The sinA are potentially useful for treating a wide range of angiogenesis-associated conditions, useful for treating a wide range of angiogenesis-associated conditions, particularly cancers, diabetic retinopathy, macular degeneration, neovascular glaucoma, arthritis, psoriasis, endometriosis, angiofibroma, and polycystic kidney disease. The sinA may also be useful for diagnosis, and polycystic kidney disease. The sinA may also be useful for diagnosis, dingle-nucleotide polymorphisms). The present sequence is used in the exemplification of the present invention.
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                                                                                                                                                                                                                                         nucleic acid (siMa) that downregulates expression of the vascular endothelial growth factor receptor (VEGRR) gene. Also described: (1) a siNA that downregulates the VEGR gene; (2) kits for in vitro or or in vitro or 
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                                                      double-stranded interfering nucleic acid, useful e.g. for treatment diagnosis of cancer, downregulates the vascular endothelial growth
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                                                                                                                                                                                                                           present invention describes a double-stranded short interfering
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                                                                                                                                                                  Example 3; SEQ ID NO 2149; 207pp; English.
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2002US-0399348P.
2002US-0406784P.
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2002US-0363124P.
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                                                                                                               factor receptor gene.
WPI; 2003-679876/64
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29-JUL-2002;
29-AUG-2002;
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06-JUN-2002;
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ADF37895
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The present invention describes a double-stranded short interfering nucleic acid (siNA) that downregulates expression of the vascular endothelial growth factor receptor (VEGFR) gene. Also described: (1) a siNA that downregulates the VEGF gene; (2) kits for in vitro or in vivo delivery of siNA; (3) conjugates and/or complexes of siNA; (4) vectors that express siNA, and (5) single-stranded siNA with similar properties. The siNAs have antianglogenic, cytostatic, antidiabetic, ophthalmological, antiarthritic, antipsociatic, nephratrocropic and gynaecological activities. The siNA are useful for modulating (downregulating) the expression of VEGFR genes. The siNA are potentially useful for treating a wide range of angiogenesis-sascoiated conditions, particularly cancers, diabetic retinopathy, macular degeneration, neovascular glaucoma, arthritis, psoriasis, endometriosis, angiofibroma, and polycystic kidney disease. The siNA may also be useful for diagnosis, and polycystic kidney disease. The siNA may also be useful for diagnosis, engineering, studying gene function, and also for gene mapping (e.g. of single-incleant) studying gene function, and also for gene mapping (e.g. of single-incleant).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     New double-stranded interfering nucleic acid, useful e.g. for treatment and diagnosis of cancer, downregulates the vascular endothelial growth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           90.5%; Score 19; DB 10; Length 21; 100.0%; Pred. No. 1.8; 0; Indels ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 21 BP; 6 A; 5 C; 4 G; 2 T; 4 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       double-stranded short interfering nucleic acid;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 3; SEQ ID NO 2092; 207pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  exemplification of the present invention
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2002US-0408378P.
2002US-0409293P.
2002US-00287949.
2002US-00306747.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BP.
                                                                                                                                                                                                                                                                                                         dcswiggen J, Beigelman L,
                                                                                                                                                                                                                              (RIBO-) RIBOZYME PHARM INC.
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ID ADF37739 standard; RNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        factor receptor gene.
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                                                                          04-NOV-2002;
27-NOV-2002;
                                                                                                                                                    15-JAN-2003;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      12-FEB-2004
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Unidentified.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADG29884/
 ò
                                                                                                                                                                                                                                                                                                                                                           The present invention describes a double-stranded short interfering nucleic acid (siNA) that downregulates expression of the vascular endothelial growth factor receptor (VEGFN) gene. Also described: (1) a siNA. that downregulates the VEGF gene. Also described: (1) a calivery of siNA; (3) conjugates and/or complexes of siNA; (4) vectors that express siNA; and (5) single-stranded siNA with similar properties. The siNAs have antianglogenic, Cytostatic, antidiabetic, ophthalmological, antiarthritic, antipsoriatic, nephrotropic and gynaecological activities. The siNA are useful for modulating (downregulating) the expression of VEGFN genes. The siNA are potentially useful for treating a wide range of angiogenesis-associated conditions, particularly cancers, diabetic retinopathy, macular degeneration, neovascular glaucoma, arthritis, postiasis, endometriosis, angiofibroma, and polycystic kidney disease. The siNA may also be useful for diagnosis, diabetic angiogenesis, and polycystic kidney disease. The siNA may also be useful for diagnosis, engineering, studying gene function, and also for gene mapping (e.g. of single-incleotide polymorphisms). The present sequence is used in the
                                                                                                                                                                                                                                                                                  New double-stranded interfering nucleic acid, useful e.g. for treatment and diagnosis of cancer, downregulates the vascular endothelial growth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         double-stranded short interfering nucleic acid; siNA; antiparkinsonian; antiarteriosclerotic; neuroprotective; nootropic; antiparkinsonian; anticonvulsant; pulmonary disease; restenoseis; atherosclerosis; Alzheimer's; Parkinson's; epilepsy; dementia; huntingcon's; amyotrophic lateral sclerosis; gene therapy; ss; DNA-RNA hybrid; FLTI.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      90.5%; Score 19; DB 10; Length 21; 78.9%; Pred. No. 1.8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 21 BP; 4 A; 4 C; 5 G; 2 T; 6 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FLT1-targeted sinA DNA-RNA hybrid - SEQ ID 402.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                       Example 3; SEQ ID NO 2152; 207pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        exemplification of the present invention.
                                                                                                                                                                                                                                  Pavco P;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 CUGAGUUUAAAAGGCACCC 19
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ADG29836/c
ID ADG29836 standard; RNA; 21 BP.
                                                                        2002US-0386782P
2002US-0393796P
                                                                                                  29-JUL-2002; 2002US-0399348P.
29-AUG-2002; 2002US-0406784P.
                                                                                                                                                     2002US-00287949.
                                                               2002WO-US017674
                                                                                                                            2002US-0408378P
                                                                                                                                         2002US-0409293P
            20-FEB-2003; 2003WO-US005022
                                                   2002US-0363124P
                                                                                                                                                                               2003US-0440129P
                                                                                                                                                                                                                                  Mcswiggen J, Beigelman L,
                                                                                                                                                                                                        (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Ouery Match
Best Local Similarity 78.3-
Best Local Similarity 78.3-
Local Similarity 78.3-
                                                                                                                                                                                                                                                                                                                factor receptor gene.
                                                                                                                                                                                                                                                          WPI; 2003-679876/64.
                                                            29-MAY-2002;
06-JUN-2002;
03-JUL-2002;
                                                                                                                                         09-SEP-2002;
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The invention relates to a double-stranded short interfering nucleic acid (sinA) molecule that down-regulates expression of an endogenous mammalian target gene comprising one or more chemical modifications and each strand of the double-stranded sinA comprises about 21 nucleotides. The sinA of the invention demonstrates antiarteriosclerotic, neuroprotective, nootropic, antiparkinsonian and anticonvulsant activities and may be useful for down-regulating the expression of an endogenous mammalian target gene and therefore in the treatment of any disease or condition that responds to modulation of gene expression or activity in a cell, tissue or organism. The disease or condition may include pulmonary diseases such as restenosis, atherosclerosis, Alzheimer's disease, pellapsy, dementa, huntington's disease, pellapsy, dementa, huntington's disease or amyotrophic lateral sclerosis. Furthermore, the sinA may be utilised for gene, therapy, applications. The current sequence is that of the sinA DNA-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New double-stranded short interfering nucleic acid molecule, useful for down-regulating the expression of an endogenous mammalian target gene or for treating diseases that respond to modulation of gene expression or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ..
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    X,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Fosnaugh
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 21 BP; 4 A; 4 C; 5 G; 2 T; 6 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FLT1-targeted sinA DNA-RNA hybrid - SEQ ID 450.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 24; SEQ ID NO 402; 593pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Beigelman L, Chowr:
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                                                                                                                                                                                                                                                           20.FEB-2002; 2002US-0358580P.
11-MAX-2002; 2002US-0363124P.
06-UIN-2002; 2002US-0386782P.
29-AUG-2002; 2002US-0406784P.
05-SEP-2002; 2002US-0408378P.
15-JAN-2003; 2003US-0440129P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (SIRN-) SIRNA THERAPEUTICS INC
                                                                                                                                                                                                20-FEB-2003; 2003WO-US005028
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RNA hybrid of the invention.
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hes 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 J, Bels.
R. Usman N,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2003-731676/69.
                                                           WO2003074654-A2
                                                                                                                             12-SEP-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Jamison S,
Synthetic.
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The invention relates to a double-stranded short interfering nucleic acid (siNA) molecule that down-regulates expression of an endogenous mammalian crarget gene comprising one or more chemical modifications and each strand of the double-stranded siNA comprises about 21 nucleotides. The siNA of the invention demonstrates antiarteriosclerotic, neuroprotective, contigories antiparkinsonian and anticonvulsant activities and may be useful for down-regulating the expression of an endogenous mammalian target gene and therefore in the treatment of any disease or condition that responds to modulation of gene expression or activity in a cell, tissue or organism. The disease or condition may include pulmonary diseases such as restenosis, atherosclerosis, Alzheimer's disease, epilepsy, dementia, huntington's disease or may experional acleral aclerosis. Furthermore, the siNA may be utilised for gene therapy applications. The current sequence is that of the siNA DNA-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New double-stranded short interfering nucleic acid molecule, useful for down-regulating the expression of an endogenous mammalian target gene or for treating diseases that respond to modulation of gene expression or activity.
amyotrophic lateral sclerosis; gene therapy; ss; DNA-RNA hybrid; FLT1.
                                                                                                                                                                                                                                                                                                                                                                                                                     Beigelman L, Chowrira B, Pavco P, Fosnaugh K;
Isman N, Thompson J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        90.5%; Score 19; DB 10; Length 21; 78.9%; Pred. No. 1.8;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 21 BP; 4 A; 4 C; 5 G; 2 T; 6 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 24; SEQ ID NO 450; 593pp; English.
                                                                                                                                                                                                                   20-FEB-2002; 2002US-0358580P.

11-MAR-2002; 2002US-0363124P.

06-JUN-2002; 2002US-0386782P.

29-AUG-2002; 2002US-0406784P.

05-SEP-2002; 2002US-0409378P.

09-SEP-2002; 2002US-0409378P.

15-JAN-2003; 2003US-0440129P.
                                                                                                                                                                                                                                                                                                                                                                                 (SIRN-) SIRNA THERAPEUTICS INC
                                                                                                                                                                            20-FEB-2003; 2003WO-US005028
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity 78.9
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                       Mcswiggen J, Beigelm
Jamison S, Usman N,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2003-731676/69.
                                                                                               WO2003074654-A2
                                   Unidentified.
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Search completed: August 19, 2005, 00:19:44 Job time : 418 secs 19 CTGAGTTTAAAAGGCACCC 1 g

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AR186006 Sequence
AR191755 Sequence
AR322637 Sequence
AR489901 Sequence
AR186005 Sequence
AR191754 Sequence
AR222636 Sequence
AR22635 Sequence
AR22635 Sequence
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AR32635 Sequence
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A98496 Sequence
AR053801 Sequence
AR063801 Sequence
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AR298136 Sequence
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GenCore version 5.1.6 Copyright (c) 1993 - 2005 Compugen Ltd.
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Maximum Match 100%
Listing first 45 summaries
                                                     OM nucleic - nucleic search, using sw model
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AR191754
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CQ831028
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61.0 21 6 CQ831032 61.0 29 6 BD253259 60.0 29 6 BD253259 60.0 21 6 AR305162 60.0 21 6 AR305162 60.0 21 6 AR305162 60.0 22 6 AR30949 59.0 22 6 AX189449 59.0 22 6 AX189449 59.1 20 6 AX266235 58.1 24 6 AX266235 58.1 24 6 AX26032 58.1 24 6 AX29431 58.1 26 6 AR39446 58.1 26 6 AR394910 58.1 26 6 AR394910 58.1 26 6 AR394910 58.1 26 6 AR394910 58.1 30 6 AR197413 58.1 26 6 AX29635 58.1 26 6 AX29635 58.1 26 6 AX29635 58.1 26 6 AX29635 58.1 26 6 AX291310 58.1 26 6 AX391310 58.1 26 6 AX391310 58.1 26 6 AX391310 58.1 26 6 AX394310 58.1 26 6 AX394310 58.1 26 6 AX394310 58.1 26 6 AX394310 58.1 26 6 AX394313 58.1 27 6 AX336422 57.1 28 6 AR324725	ALIGNMENTS 6 1494 from patent US 6346398. 6.1 GI:20231971 ified. 7 McSwiggen,J., Stinchcomb,D. and and reagent for the treatment of di to levels of vascular endothelial to levels of vascular endothelial Location/Qualifiers	/organism="unknown" /mol_type="unassigned DNA"	17 bp DNA 7243 from patent US 63463981 GI:20237720 fied. s 1 to 17) MCSwiggen,J., Stinchcomb,D. and
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Location/Qualifiers
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Unclassified.
Unclassified.
I (bases I to 17)
Pavco, P. M. Swiggen, J.A., Stinchcomb, D.T. and Escobedo, J.
Method and reagent for the treatment of diseases or conditions
related to levels of vascular endothelial growth factor receptor
Patent: US 6566127-A 39 20-MAY-2003;
Method and reagent for the treatment of diseases or conditions related to levels of vascular endothelial growth factor receptor Patent: US 6346398-A 7243 12-FEB-2002; Location/Qualifiers
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Bennett, C.F. and Watt, A.T.

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Patent: US 6710174-A 24 23-MAR-2004;
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/wol_type="unassigned RNA"
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Sequence 24 from patent US 6710174.
AR489901
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Sequence 39 from patent US 6566127.
AR322637
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                                                                   /organism="unknown"
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Pavco, P., McSwiggen, J., Stinchcomb, D. and Escobedo, J.
Method and reagent for the treatment of diseases or conditions related to levels of vascular endothelial growth factor receptor Patent: US 6346398-A 1493 12-FEB-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 (bases 1 to 17)
Pavco, P., McSwiggen, J., Stinchcomb, D. and Escobedo, J.
Method and reagent for the treatment of diseases or conditions related to levels of vascular endothelial growth factor receptor Patent: US 6346398-A 7342 12-FEB-2002;
Location/Qualifiers
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AR186005
Seguence 1493 from patent US 6346398.
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AR322636.1 GI:33708444
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/mol_type="unassigned DNA"
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/mol_type="unassigned DNA"
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us-10-665-951-2185.rge

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Matches

ACCESSION VERSION KEYWORDS

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BD233486 22 bp DNA linear PAT 17-JUL-2003
Novel human thrombopoietin derivative having elevated capability of
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1 (bases 1 to 17)

Parco, P., McSwiggen, J., Stinchcomb, D. and Escobedo, J.

Bathod and reagent for the treatment of diseases or conditions related to levels of vascular endothelial growth factor receptor Patent: US 6346398-A 7241 12-FEB-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 (bases 1 to 17)
Pavco, P., McSwiggen, J.A., Stinchcomb, D.T. and Escobedo, J.
Method and reagent for the treatment of diseases or conditions
related to levels of vascular endothelial growth factor receptor
Patent: US 6566127-A 37 20-MAY-2003;
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71.4%; Score 15; DB 6; Length 17; llarity 73.3%; Pred. No. 1.9e+04; Conservative 4; Mismatches 0; Indels
   Length 17;
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Pavco,P., McSwiggen,J., Stinchcomb,D. and Escobedo,J.
Method and reagent for the treatment of diseases or conditions
related to levels of vascular endothelial growth factor receptor
Patent: US 6346398-A 1492 12-FEB-2002;
                                   1 (bases 1 to 17)
Pavco,P., McSwiggen,J.A., Stinchcomb,D.T. and Escobedo,J.
Rethod and reagent for the treatment of diseases or conditions related to levels of vascular endothelial growth factor receptor Patent: US 6566127-A 38 20-MAY-2003;
Location/Qualifiers
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Bennett, C.F. and Watt, A.T.

Bennett, C.F. and Watt, A.T.

Antisense inhibition of vascular endothelial growth factor receptor-1 expression

Patent: US 6710174-A 25 23-MAR-2004;

Location/Qualifiers
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                                                                                                                                                                                                                                     76.2%; Score 16; DB 6; Length 17; 75.0%; Pred. No. 5.8e+03; ive 4; Mismatches 0; Indels
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il Similarity 81.2%; Pred. No. 5.6e+03;
13; Conservative 3; Mismarch-
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Sequence 1492 from patent US 6346398.
AR186004
                                                                                                                                                                                                                                                                                                                                                                                                                                          DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 25 from patent US 6710174.
AR489902
AR489902.1 GI:47257015
                                                                                                                                                                /organism="unknown"
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/organism="unknown"
/mol_type="genomic DNA"
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20 AGTTTAAAAGGCACCC 5
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                  Unclassified.
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Best Local Similarity
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Matches 12; Conserv
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DEFINITION ACCESSION VERSION KEYWORDS SOURCE

RESULT 9 AR186004

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LOCUS

ORGANISM

AUTHORS TITLE

JOURNAL

FEATURES

ORIGIN

REFERENCE

Gaps

; 0

linear MAM 29-JUL-1997

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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Cetartiodactyla; Suina; Suidae; Sus.

1 (bases 1 to 2)

Fridolfsson, A.K., Hori, T., Wintero, A.K., Fredholm, M., Yerle, M., Robic, A., Andersson, L. and Ellegren, H.

Robic, A., Andersson, L. and Ellegren, H.

Expansion of the pig comparative map by expressed sequence tags (EST) mapping
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Rattus sp.
Rattus sp.
Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Mammalia; Butheria; Rodentia; Sciurognathi; Muridae; Murinae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Direct Submission
Submitted (27-UUL-1997) Fridolfsson A.K., Animal Breeding and
Genetics, Swedish University of Agricultural Sciences, Biomedical
Center, Box 597, S-751 24 Uppsala, SWEDEN
Location/Qualifiers
                                                                                                 /organism='Artificial Sequence'.
                                                                                                                                                                                                                                                                                                               Score 14.2; DB 6; Length 24;
Pred. No. 4.6e+04;
3; Mismatches 3; Indels
C12P21/02,C12N15/00,A61K37/24,C12N5/00
Synthetic oligodeoxynucleotide primer 30-C
Key
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Best Local Similarity 64.7%; Pred. No. 7.7e+04;
Matches 11; Conservative 4; Mismatches 2;
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AJ000790

    .24
    forganism="synthetic construct"
|mol_type="genomic DNA"
|db_xref="taxon:32630"

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Sequence 6 from Patent WO9911784.
A98496
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/db_xref="taxon:9823"
/chromosome="12"
/map="12p"
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PCR primer.
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Fridolfsson, A.K.
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                                                                                         synthetic construct
cother sequences.

Synthetic construct
cother sequences; artificial sequences.

Chung, J.Y., Park, S.K., Ju, S.M., Kyung, H., Ahn, Lim, S.W.,
Chang, W.I., Park, S.K., Woh, Y.W. and Park, J.S.
Chung, J.Y., Park, S.K., Woh, Y.W. and Park, J.S.
Chang, W.I., Park, S.K., Woh, Y.W. and Park, J.S.
Novel human thrombopoietin derivative having elevated capability of proliferating platelets in vivo
Droliferating platelets in vivo
Dr
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S JP 2002519031-A/6.
S Synthetic construct
(SM synthetic construct
construct
other sequences; artificial sequences.
S Chung, Jr., Park, S.K., Voh, Y.W. and Park, J.S.
Chung, M.I., Park, S.K., Koh, Y.W. and Park, J.S.
Novel human thrombopoietin derivative having elevated capability of proliferating platelete in vivo
D PREWOONG PHARMACEUTICAL CO LTD
OA Artificial Sequence
PN JP 2002519031-A 6 02-JUL-2002;
D AEWOONG PHARMACEUTICAL CO LTD
OS Artificial Sequence
PN JP 2002519031-A 6 02-JUL-2002;
D AEWOONG PHARMACEUTICAL CO LTD
OS Artificial Sequence
PN JP 2002519031-A 6 02-JUL-2002;
D AEWOONG PHARMACEUTICAL CO LTD
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PN JP 2002519031-A 6 02-JUL-2002;
D AEWOONG PHARMACEUTICAL CO LTD
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Novel human thrombopoietin derivative having elevated capability of proliferating platelets in vivo.
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PR 30-JUN-1999 KR 19982514, 29-JUN-1999 KR 19992514
JOO YOUNG CHUNG, SANG KYU PARK, SANG MYOUNG JU, HYEA KYUNG PI
AHN, SEUNG WOOK LIM,
PI WOO IK CHANG, SEUNG KOOK PARK, YEO WOOK KOH, JI SOO PARK
C12N15/09, A61K38/22, A61P7/00, A61P7/04, A61P43/00, A61P43/00,
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Location/Qualifiers
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    .22
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  proliferating platelets in vivo.
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                        BD233486
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JP 2002519031-A/5.
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Gaps

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Indels

Length 21;

PAT 26-JAN-2000

linear

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Rattus.

(bases 1 to 28)

England, S. and Chen, C.

PROTON GATED ION CHANNEL PROTEINS

Patent: WO 9911784-A 6 11-MAR-1999;

ENGLAND STEVE (GB); UNIV LONDON (GB)

Location/Qualifiers

1. 28

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                                                                                                 FEATURES
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AUTHORS
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Gaps . 0 Query Match 65.7%; Score 13.8; DB 6; Length 28; Best Local Similarity 76.5%; Pred. No. 7.2e+04; Matches 13; Conservative 2; Mismatches 2; Indels 3 GAGUUDAAAGGCACCC 19 ||| ::||||||| |:|| 18 GAGGTTAAAAGGCTCCC 2 a

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